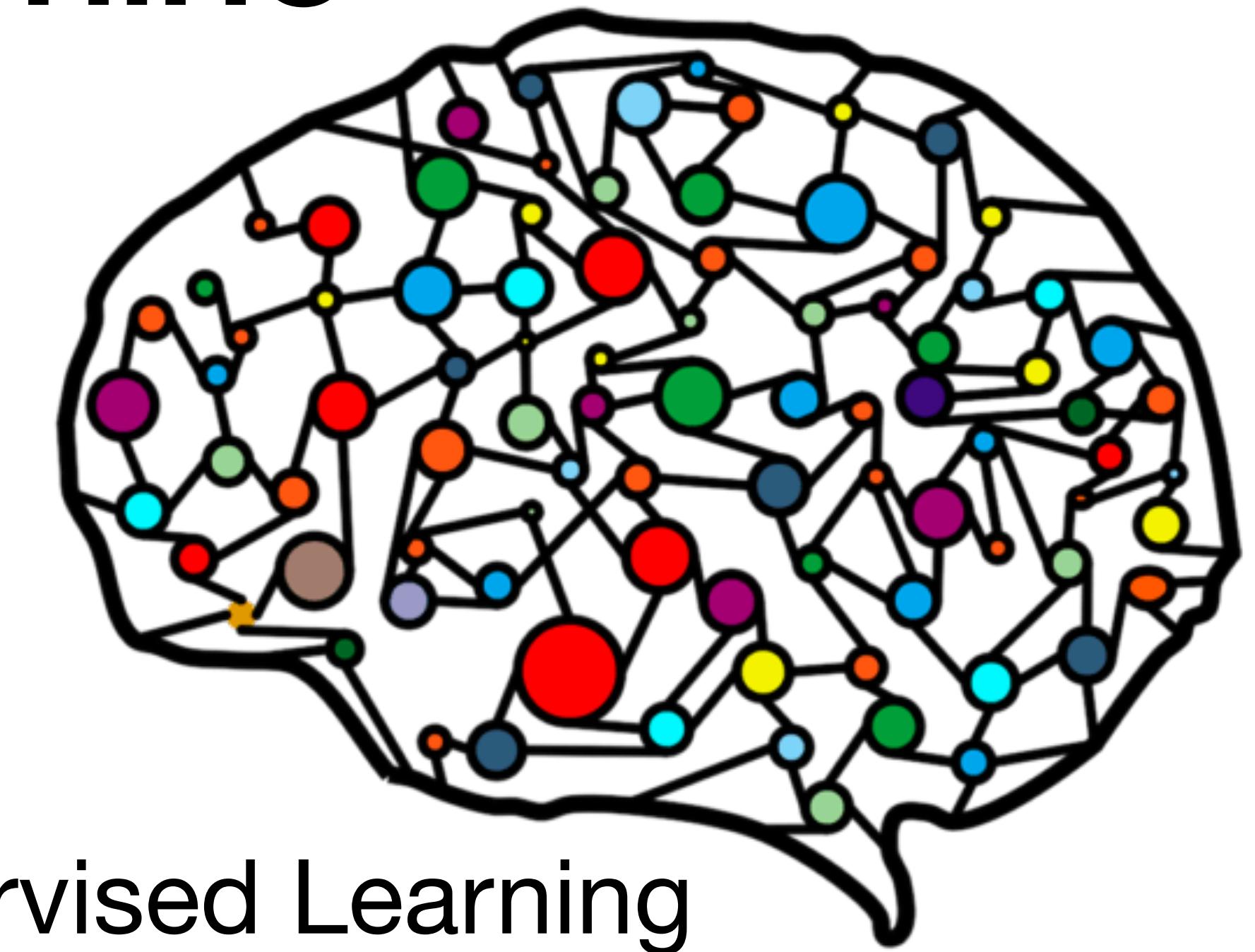


General Principles of Human and Machine Learning



Lecture 9: Supervised and Unsupervised Learning

Dr. Charley Wu

<https://hmc-lab.com/GPHML.html>

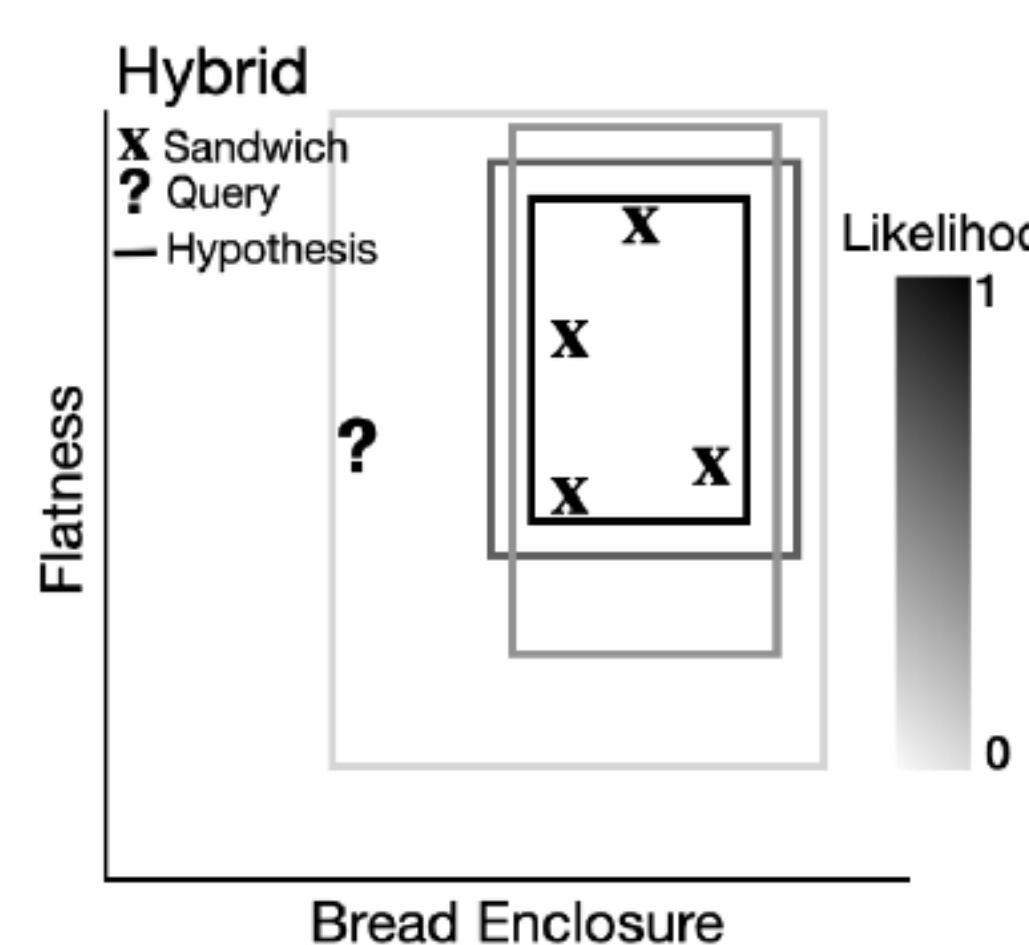
Any clarification questions from
previous weeks?

Teaching evaluations

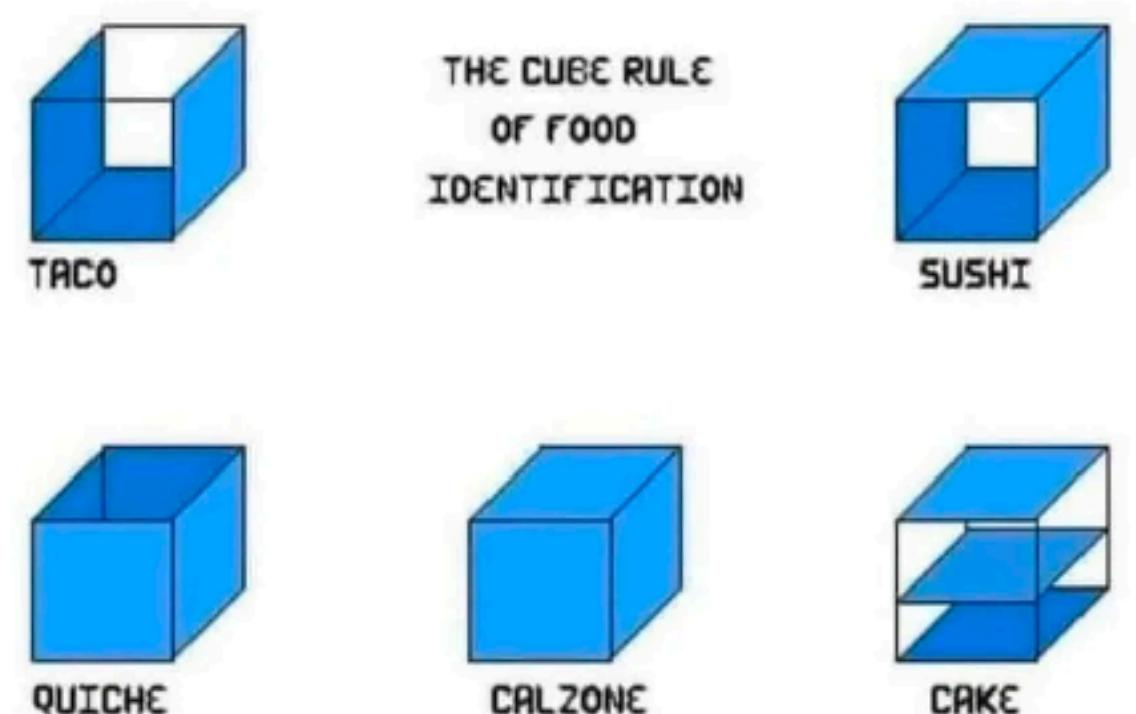
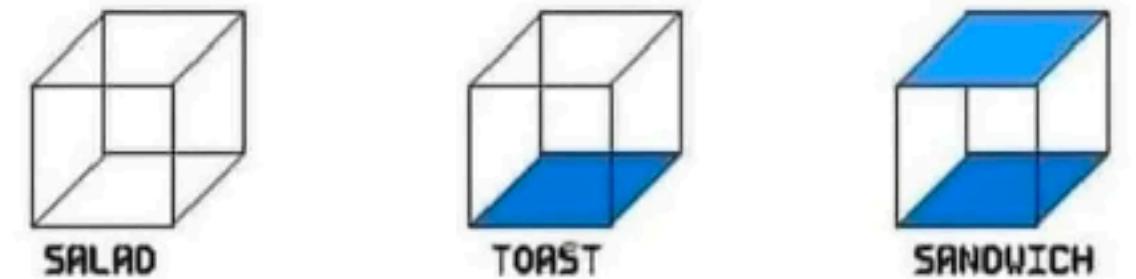
- You will receive an email to submit your evaluations of this course between 06.01. to 20.01
- Your feedback will be very helpful for us, particularly Alex and David who quite new to teaching
- None of us are paid to teach, and we are organizing this course entirely from our own interest
- This will also be my last class at Uni Tübingen

Last week ...

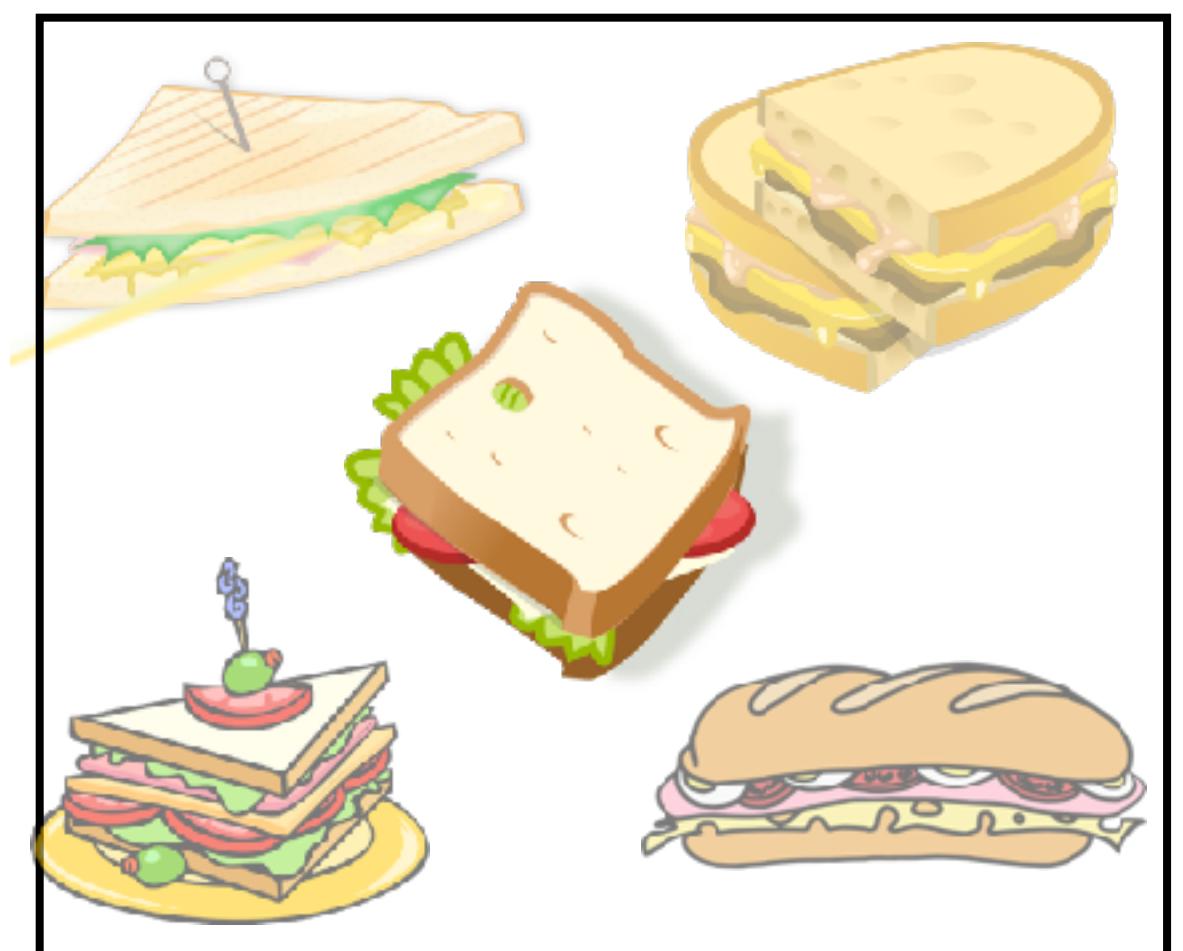
- Concepts are mental representations of categories in the world
- Classical view used **rules** to describe the necessary and sufficient conditions for category membership
- More psychological approaches used **similarity**, compared to learned prototypes or past exemplars
- Bayesian concept learning is a **hybrid** approach, that uses distributions over rules, and recreating patterns consistent with similarity-based approaches



Rules



Similarity



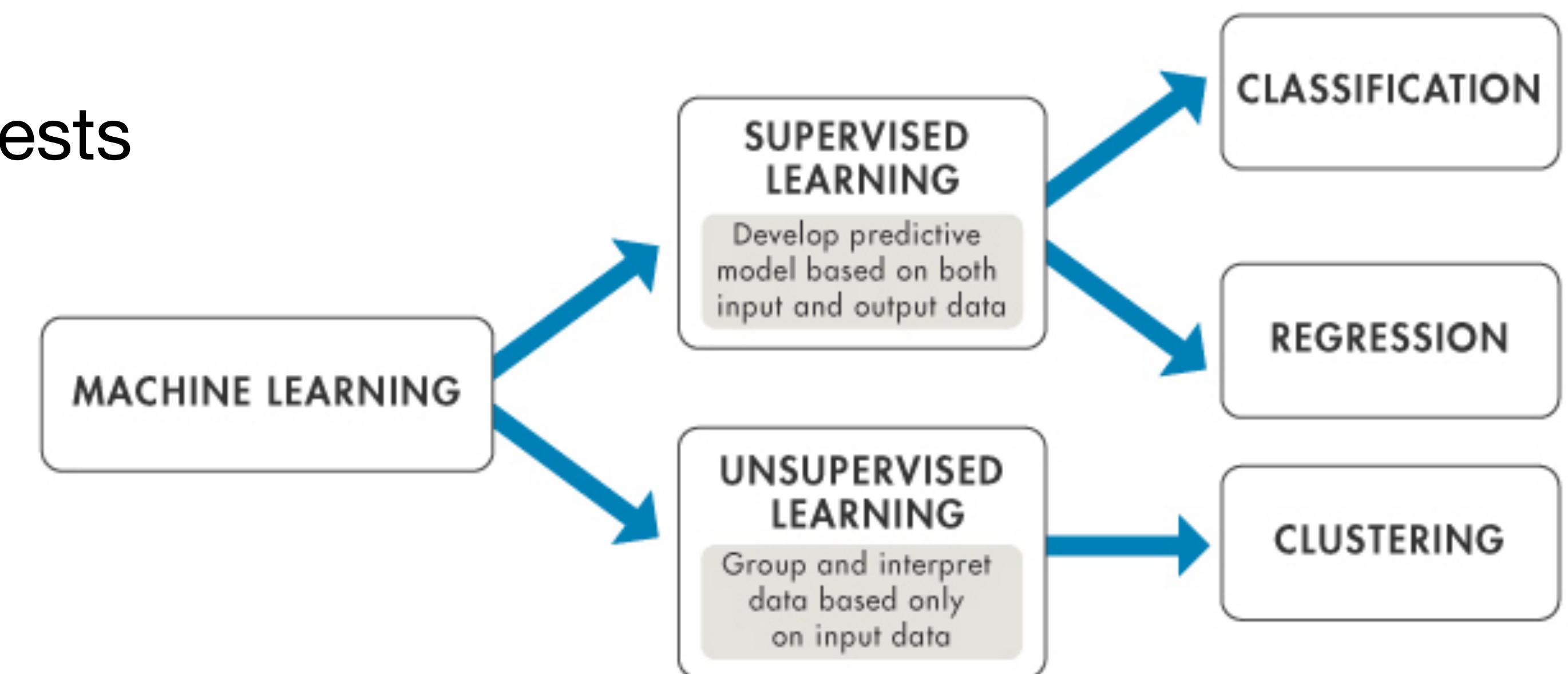
Today's agenda

Supervised learning (for classification)

- Multilayer Perceptrons
- Decision trees and random forests
- Support vector machines
- Naïve Bayes

Unsupervised Learning

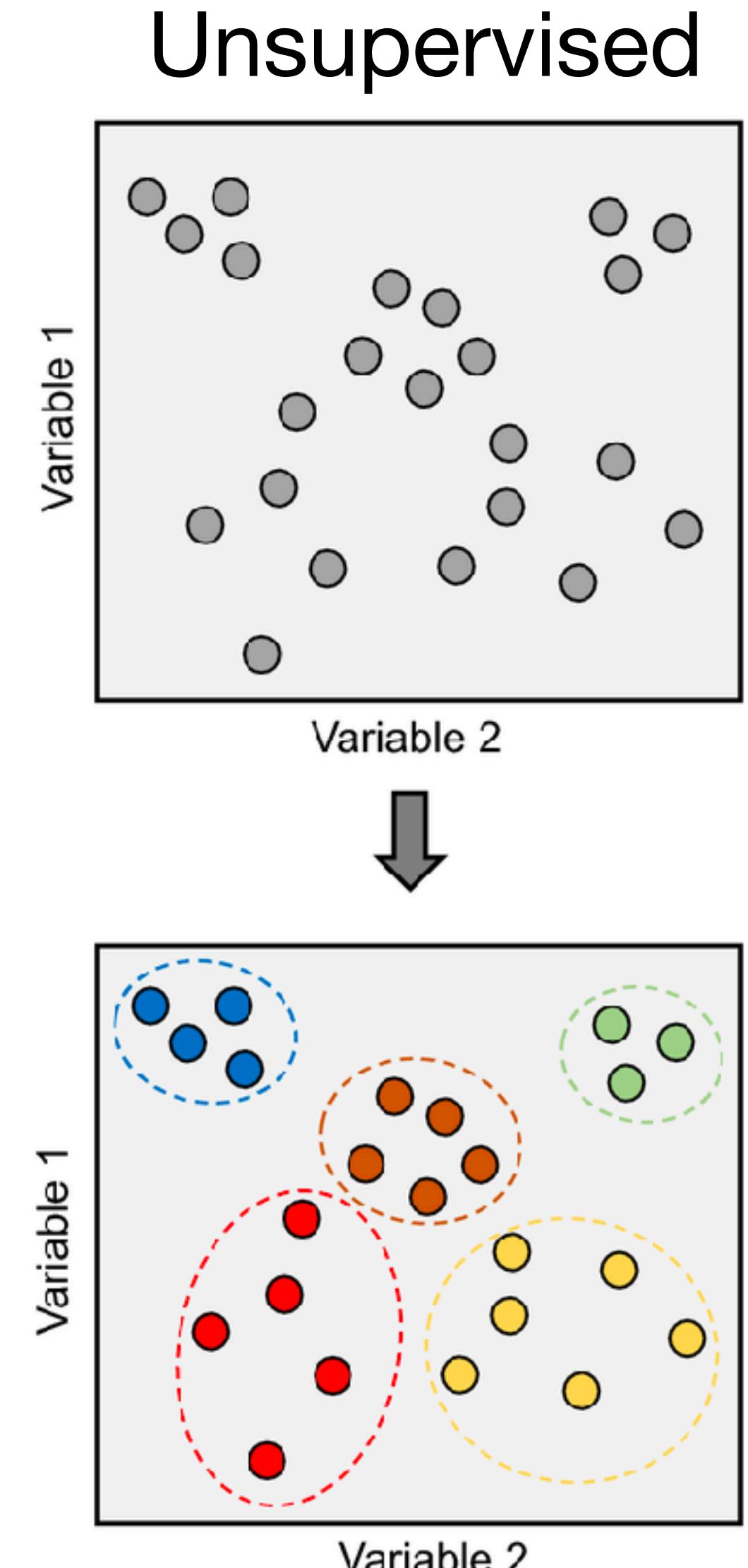
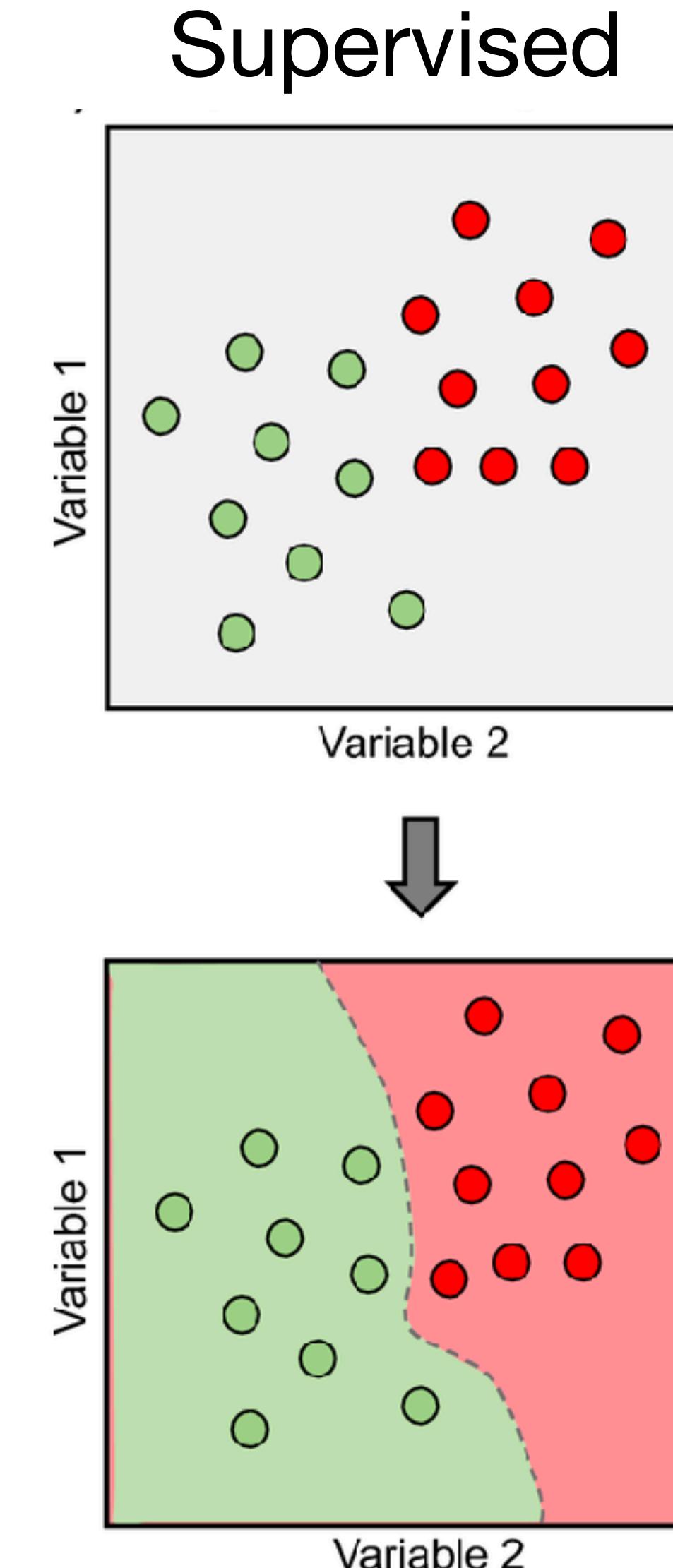
- k-Means
- Gaussian Mixture models



Supervised vs. unsupervised learning

- Classification problems*: classify data points into one of n different categories
- **Supervised** learning:
 - Training data provides category labels
 - Classifiers usually try to learn a decision-boundary
- **Unsupervised** learning:
 - Training data lacks category labels
 - Classifiers usually try to learn clusters

*Note that *regression* is another class of ML problems, which we will discuss next week



Supervised learning

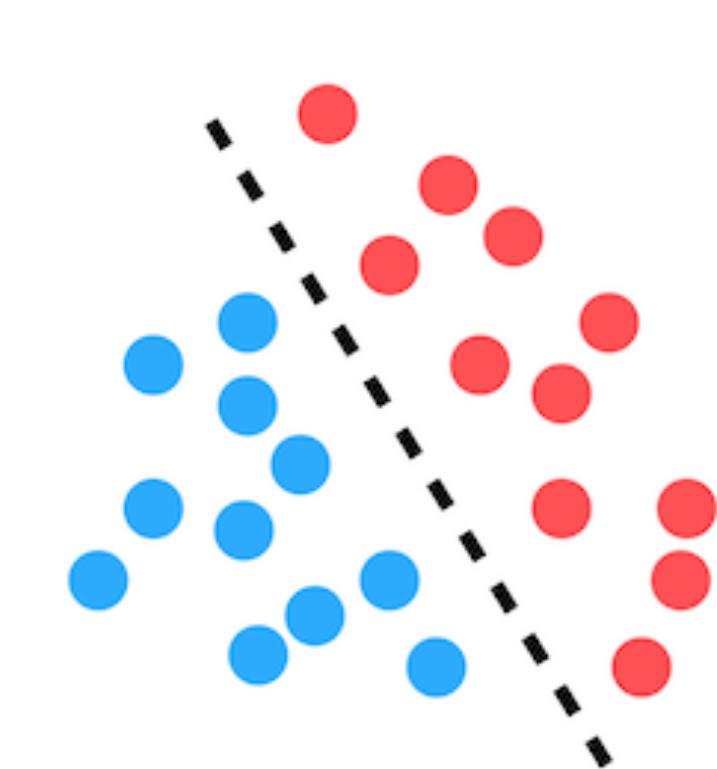
- Two general classes:
 - **Discriminitive** directly map features to class labels, often by learning a decision-boundary (rule-like)
 - **Generative** approaches learn the probability distribution of the data (similarity-like)
- Example problem: Spam detector
 - Data $\mathcal{D} = \{\mathbf{X}, \mathbf{y}\}$
 - each $\mathbf{x} \in \mathbf{X}$ are the features of an email (e.g., length, date, sender, content, etc...)
 - each $y \in \mathbf{y}$ is the label (1 if spam, 0 otherwise)
- **Discrimitive** models identify the boundaries that separate spam from non-spam
- **Generative** models learn the distributions of spam and non-spam emails

Notation:

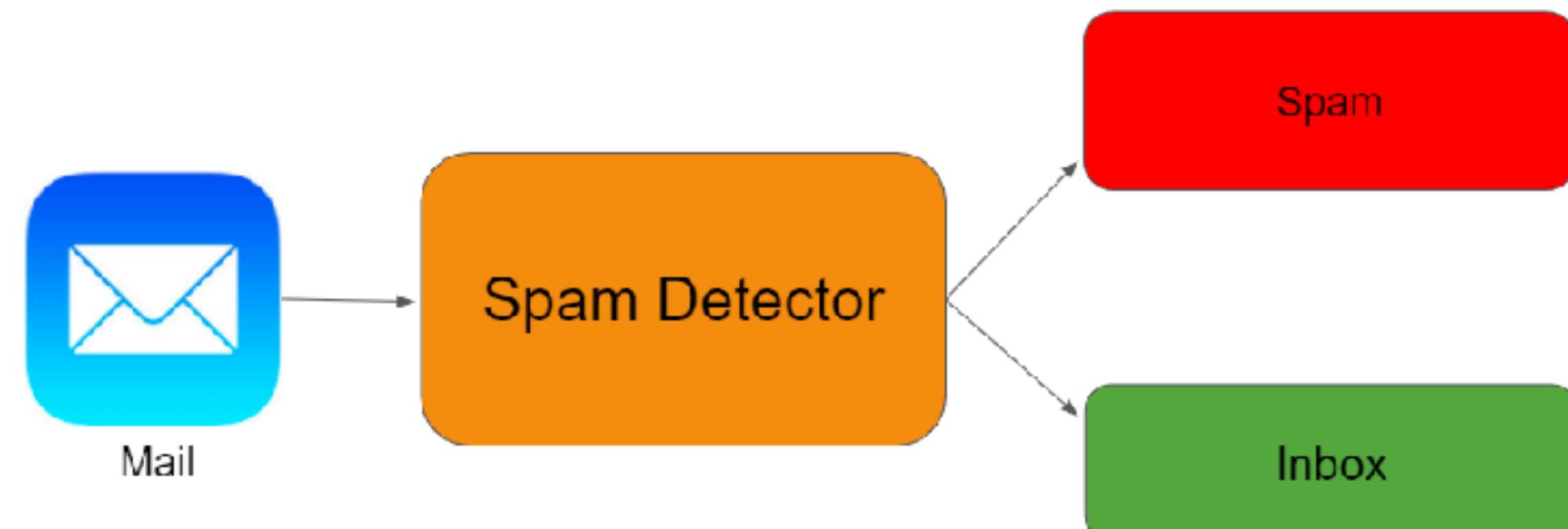
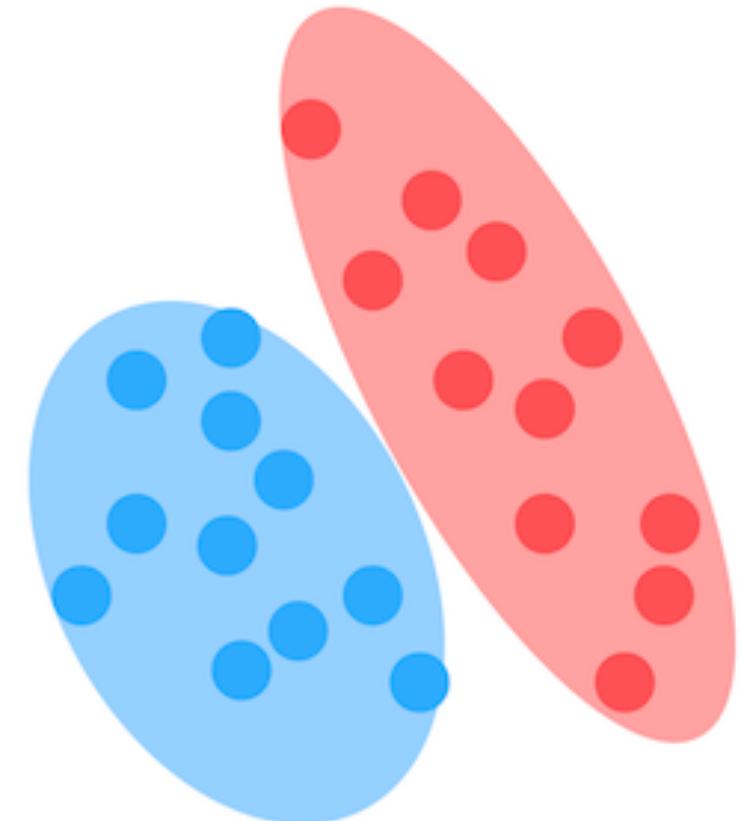
a scalar
 A constant

\mathbf{a} vector
 \mathbf{A} Matrix

Discriminative



Generative



Perceptrons and Neural Networks

- Perceptrons were the first ML classifiers
- More generally, Multilayer Perceptrons (MLPs) can learn any arbitrary decision boundary (i.e., non-linear) by adding more hidden layers
- Universal approximation theorem (Cybenko, 1989)
- Training via backpropagation

MSE Loss

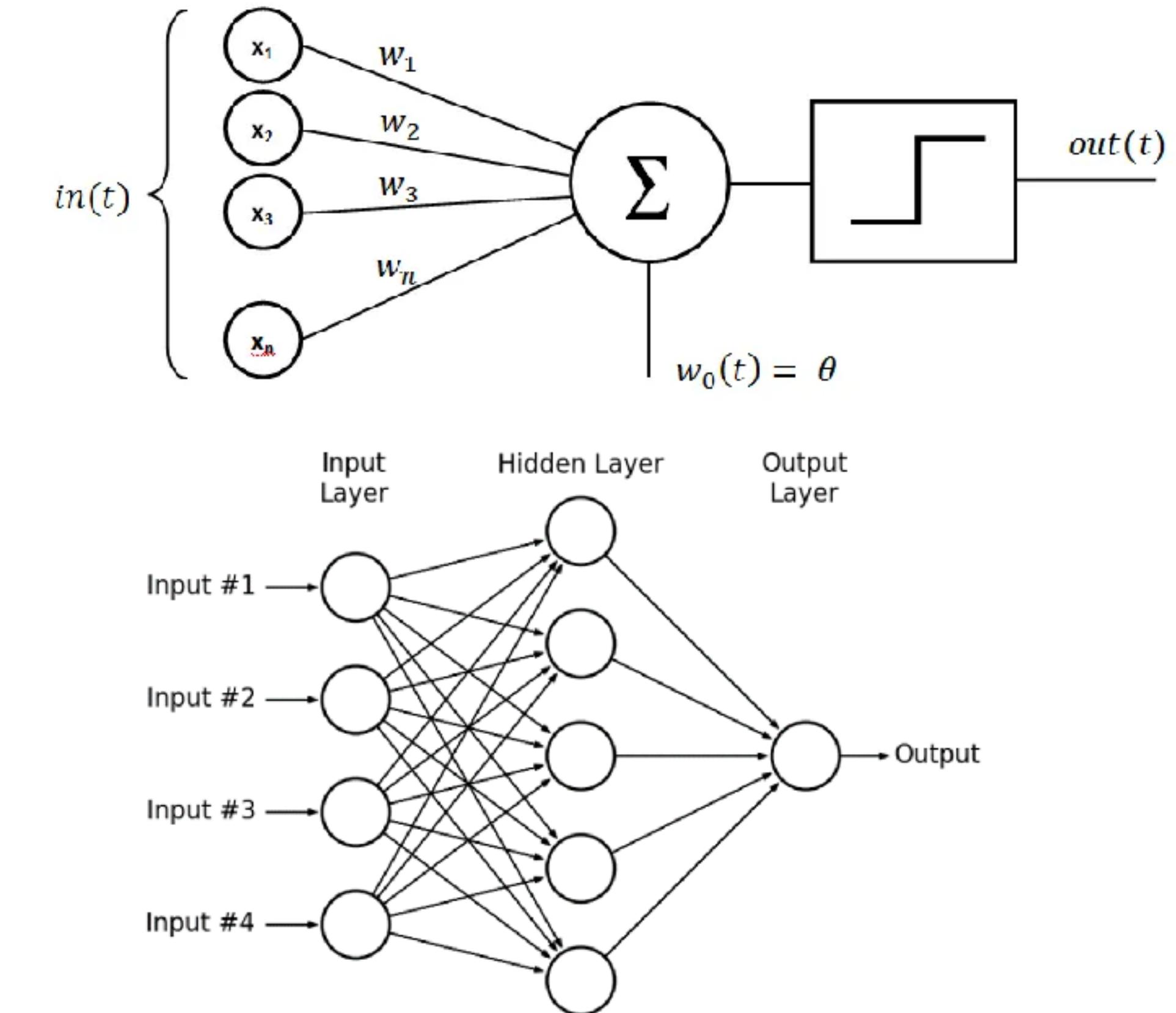
$$\mathcal{L} = \frac{1}{2} \sum_{i=1}^m (y_i - \hat{y}_i)^2$$

Weight updates

$$w \leftarrow w - \alpha \frac{\partial \mathcal{L}}{\partial w}$$

where

$$\frac{\partial \mathcal{L}}{\partial \mathbf{w}} = \frac{\partial \mathcal{L}}{\partial \mathbf{u}} \frac{\partial \mathbf{u}}{\partial \mathbf{w}}$$



Perceptrons and Neural Networks

- Perceptrons were the first ML classifiers
- More generally, Multilayer Perceptrons (MLPs) can learn any arbitrary decision boundary (i.e., non-linear) by adding more hidden layers
- Universal approximation theorem (Cybenko, 1989)
- Training via backpropagation

MSE Loss

$$\mathcal{L} = \frac{1}{2} \sum_{i=1}^m (y_i - \hat{y}_i)^2$$

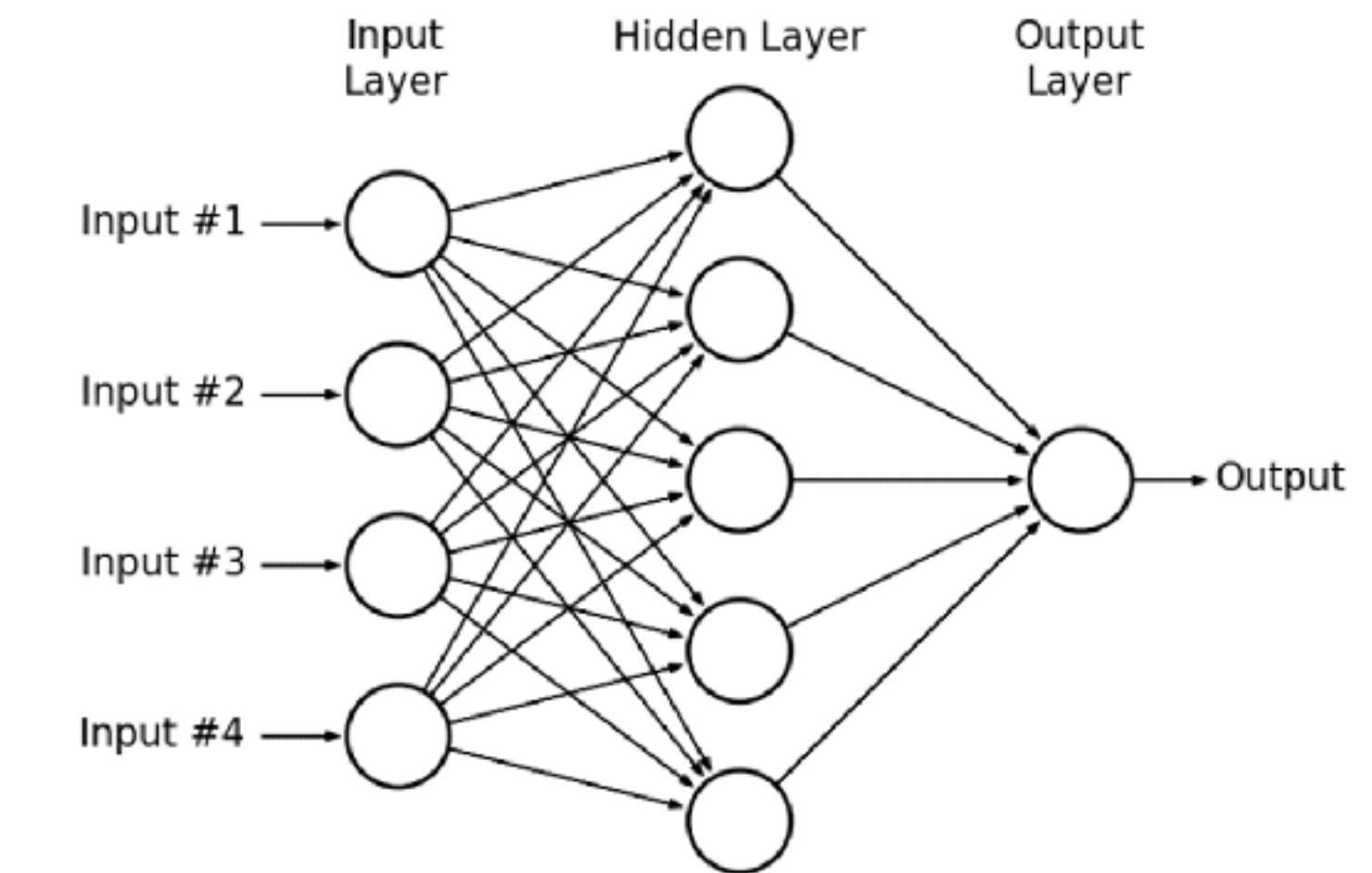
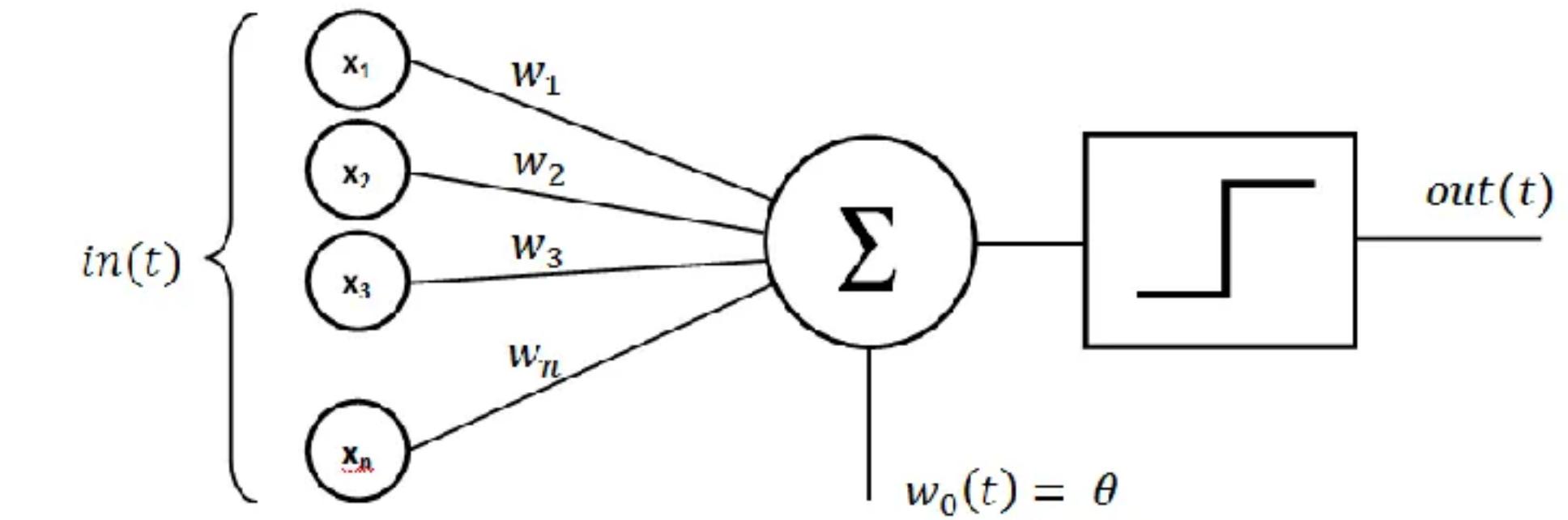
↑
prediction

Weight updates

$$w \leftarrow w - \alpha \frac{\partial \mathcal{L}}{\partial w}$$

where

$$\frac{\partial \mathcal{L}}{\partial \mathbf{w}} = \frac{\partial \mathcal{L}}{\partial \mathbf{u}} \frac{\partial \mathbf{u}}{\partial \mathbf{w}}$$



Perceptrons and Neural Networks

- Perceptrons were the first ML classifiers
- More generally, Multilayer Perceptrons (MLPs) can learn any arbitrary decision boundary (i.e., non-linear) by adding more hidden layers
- Universal approximation theorem (Cybenko, 1989)
- Training via backpropagation

MSE Loss

$$\mathcal{L} = \frac{1}{2} \sum_{i=1}^m (y_i - \hat{y}_i)^2$$

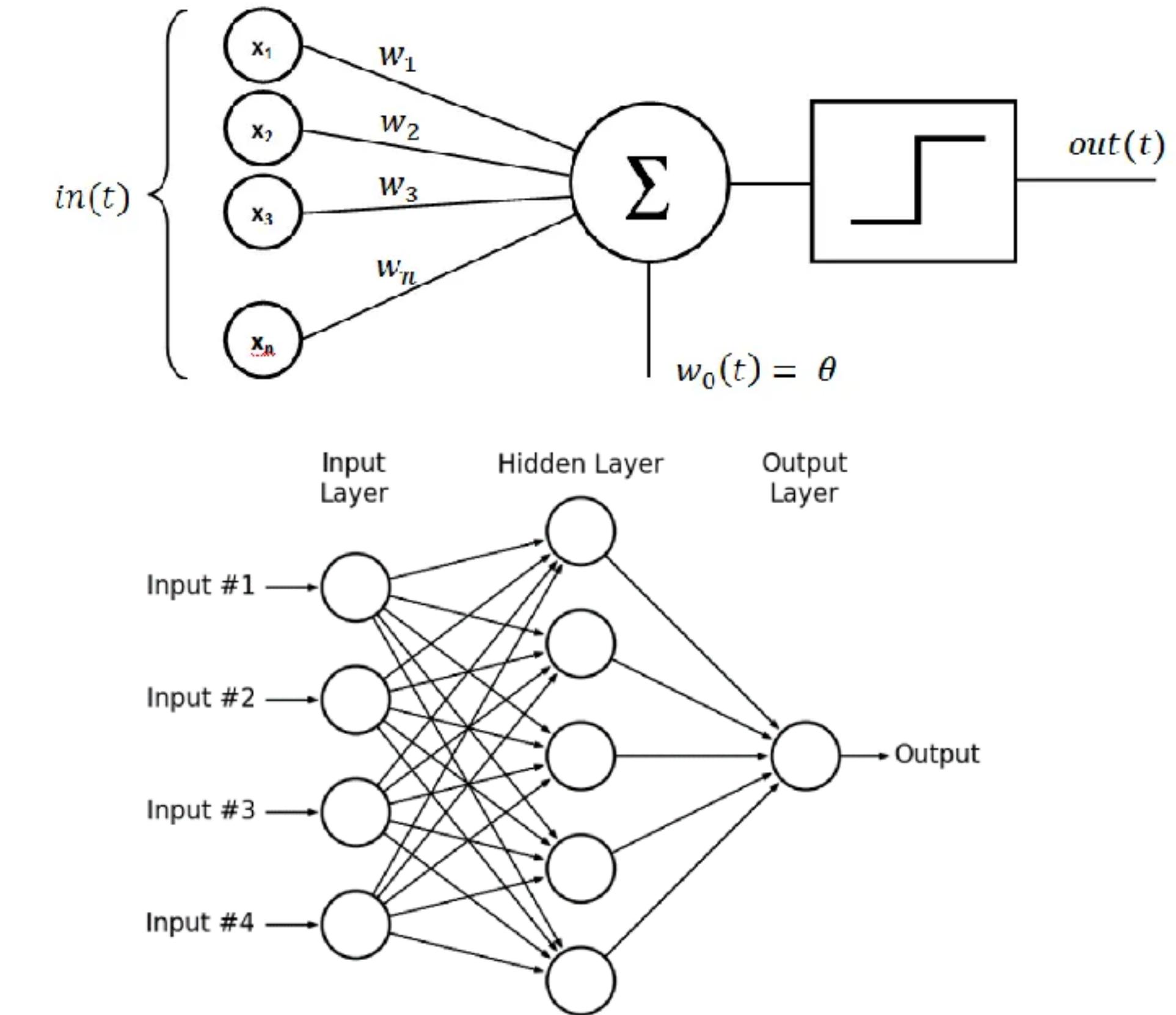
↑
prediction

Weight updates

$$w \leftarrow w - \alpha \frac{\partial \mathcal{L}}{\partial w}$$

↑
learning rate

where $\frac{\partial \mathcal{L}}{\partial \mathbf{w}} = \frac{\partial \mathcal{L}}{\partial \mathbf{u}} \frac{\partial \mathbf{u}}{\partial \mathbf{w}}$



Perceptrons and Neural Networks

- Perceptrons were the first ML classifiers
- More generally, Multilayer Perceptrons (MLPs) can learn any arbitrary decision boundary (i.e., non-linear) by adding more hidden layers
- Universal approximation theorem (Cybenko, 1989)
- Training via backpropagation

MSE Loss

$$\mathcal{L} = \frac{1}{2} \sum_{i=1}^m (y_i - \hat{y}_i)^2$$

↑
prediction

Weight updates

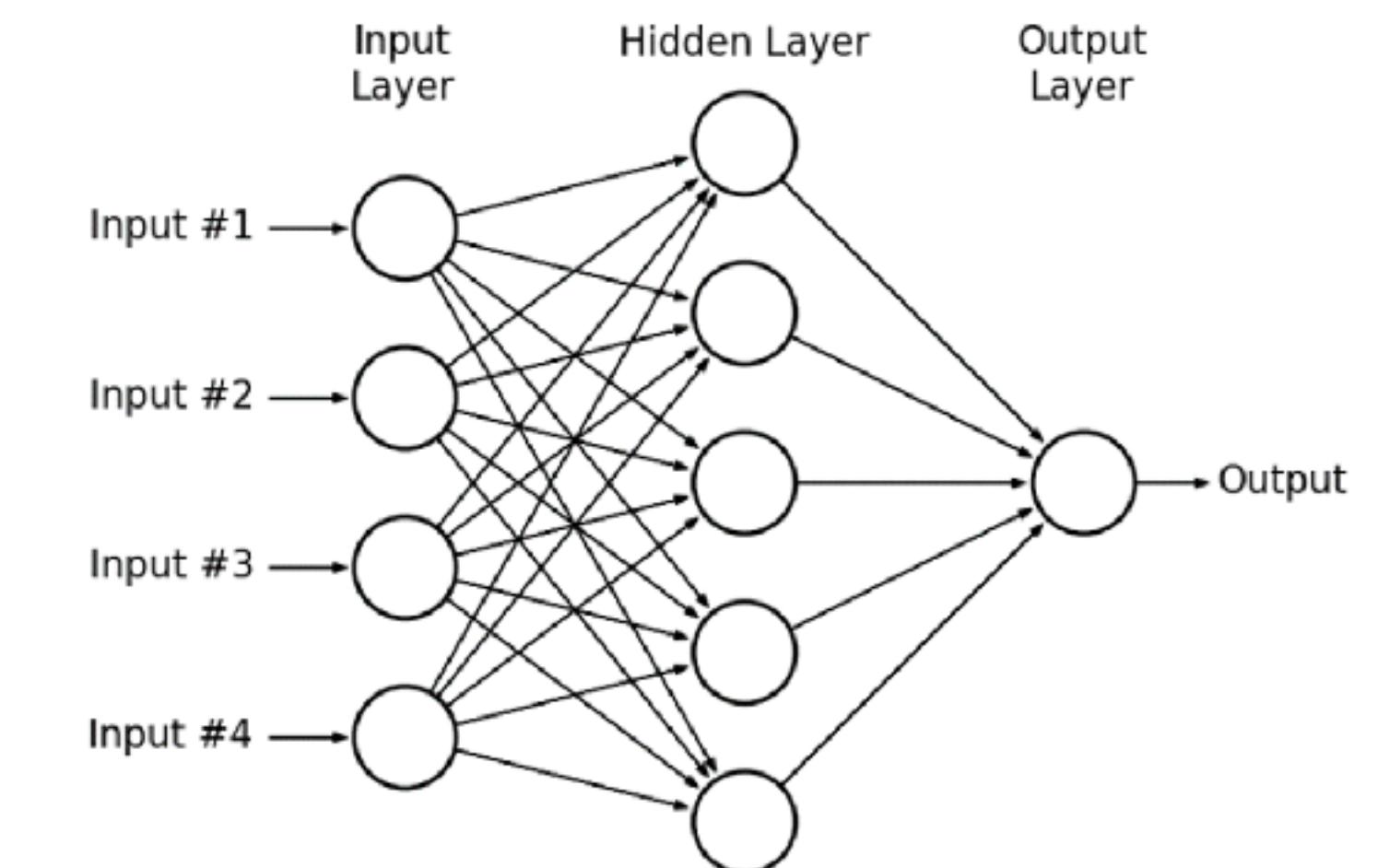
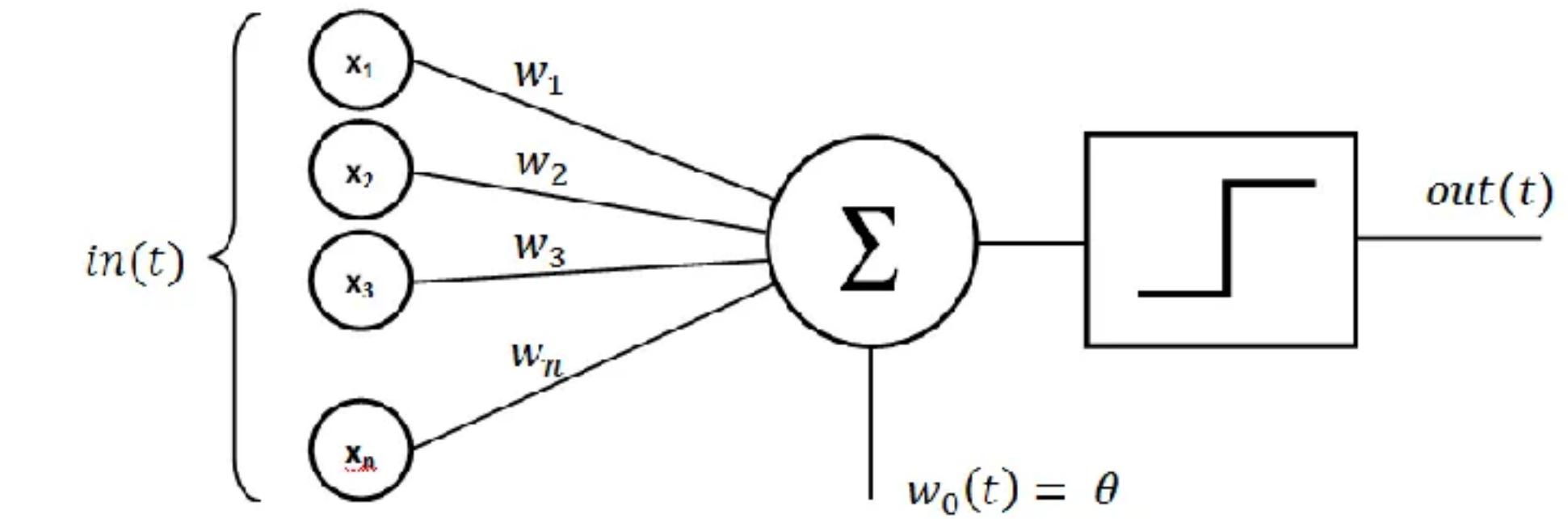
$$w \leftarrow w - \alpha \frac{\partial \mathcal{L}}{\partial w}$$

↑
learning rate

where

$$\frac{\partial \mathcal{L}}{\partial \mathbf{w}} = \frac{\partial \mathcal{L}}{\partial \mathbf{u}} \frac{\partial \mathbf{u}}{\partial \mathbf{w}}$$

Chain rule is used to pass derivatives over layers



Perceptrons and Neural Networks

- Perceptrons were the first ML classifiers
- More generally, Multilayer Perceptrons (MLPs) can learn any arbitrary decision boundary (i.e., non-linear) by adding more hidden layers
- Universal approximation theorem (Cybenko, 1989)
- Training via backpropagation

MSE Loss

$$\mathcal{L} = \frac{1}{2} \sum_{i=1}^m (y_i - \hat{y}_i)^2$$

↑
prediction

Weight updates

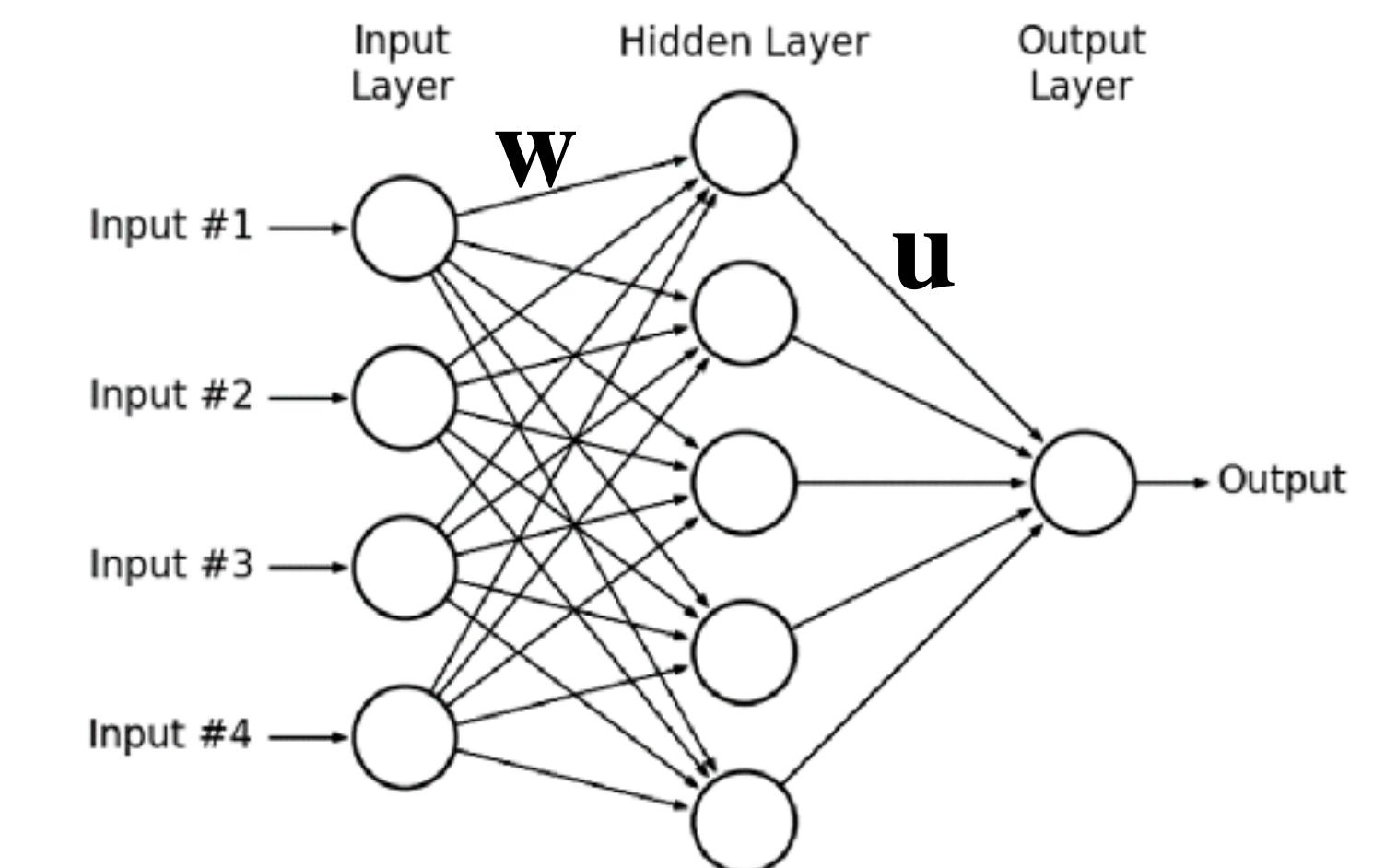
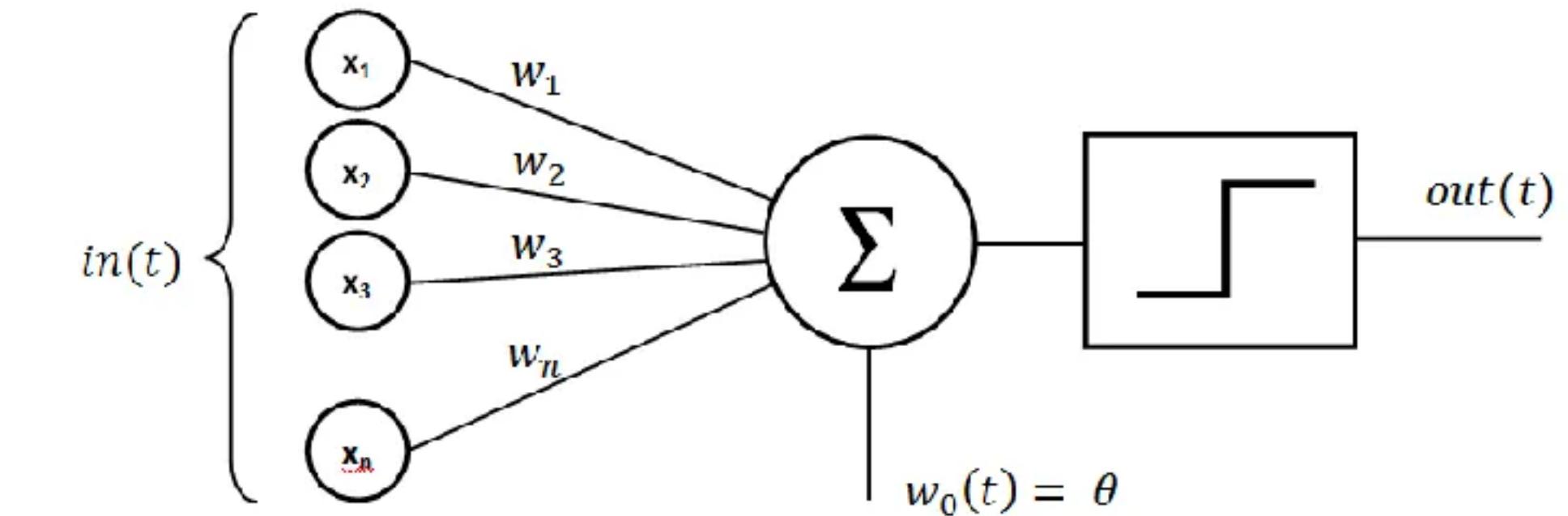
$$w \leftarrow w - \alpha \frac{\partial \mathcal{L}}{\partial w}$$

↑
learning rate

where

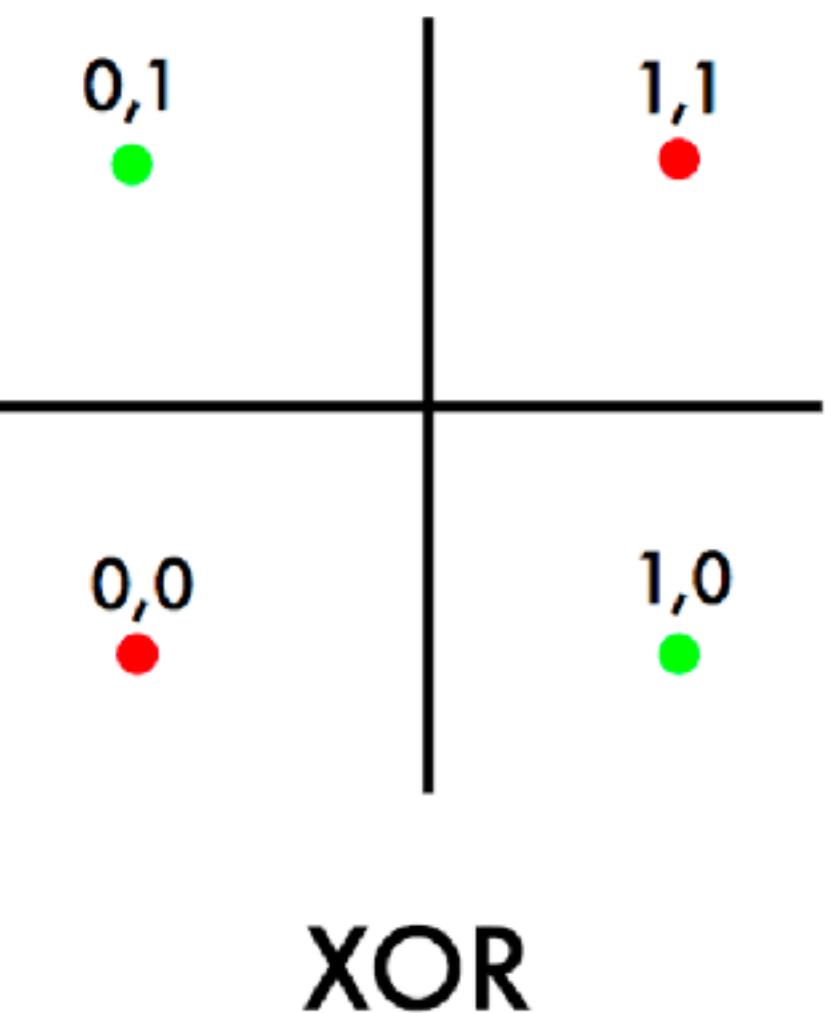
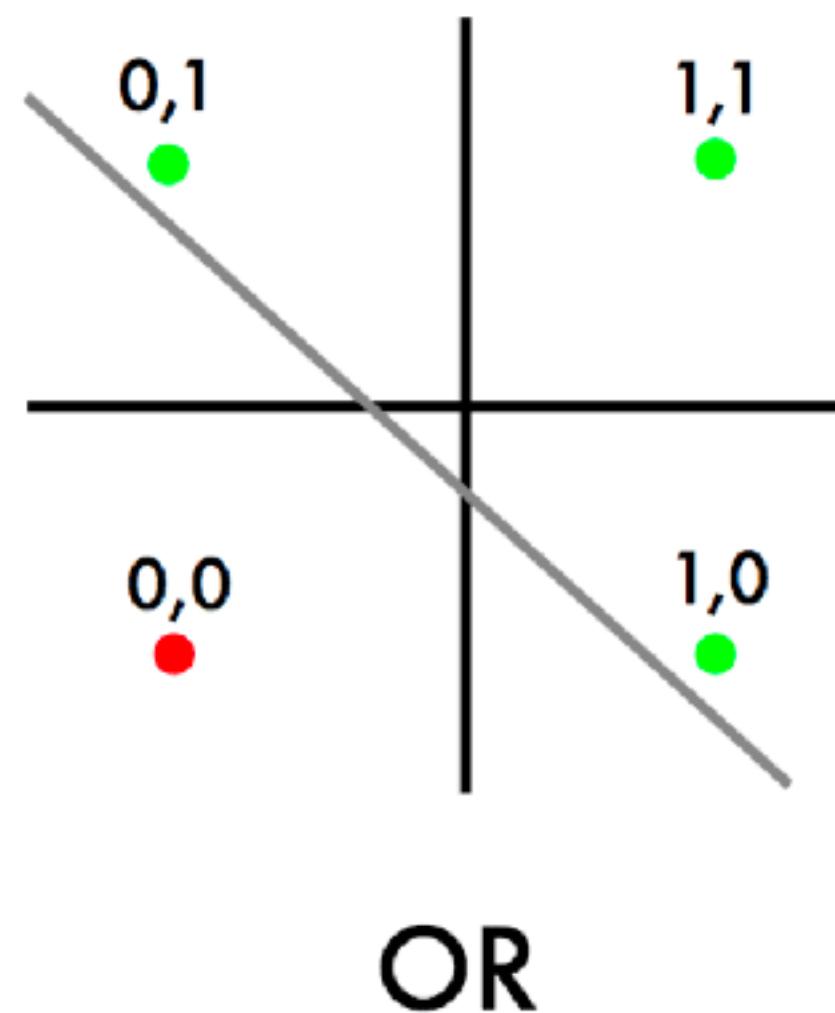
$$\frac{\partial \mathcal{L}}{\partial \mathbf{w}} = \frac{\partial \mathcal{L}}{\partial \mathbf{u}} \frac{\partial \mathbf{u}}{\partial \mathbf{w}}$$

Chain rule is used to pass derivatives over layers

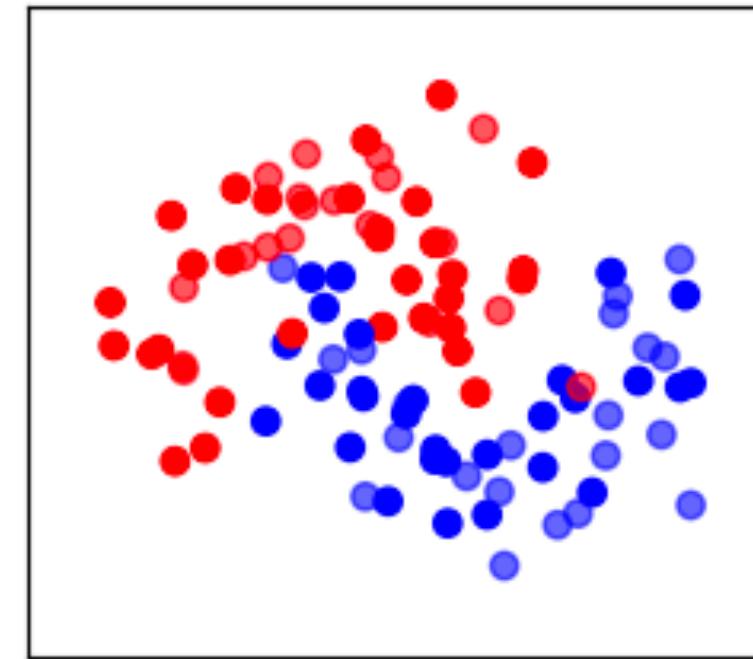


Decision boundaries

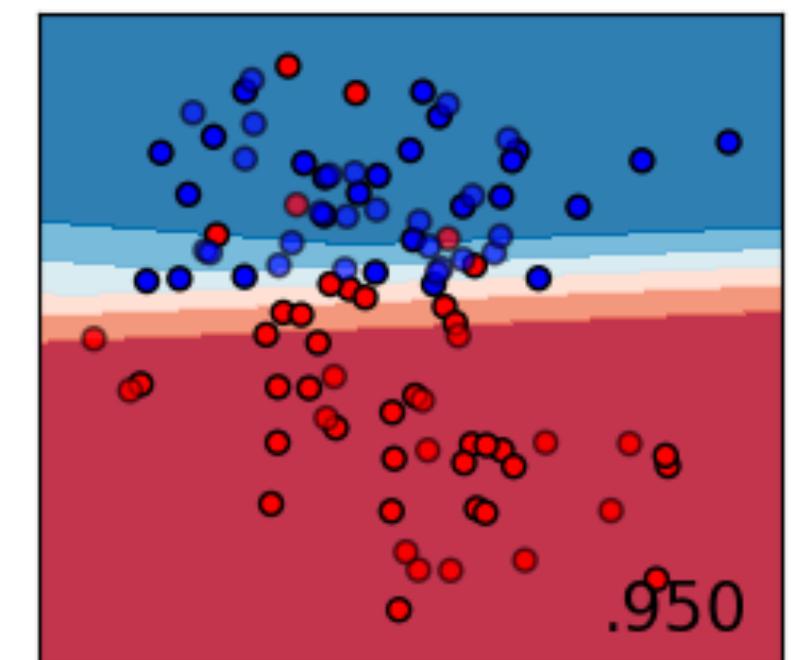
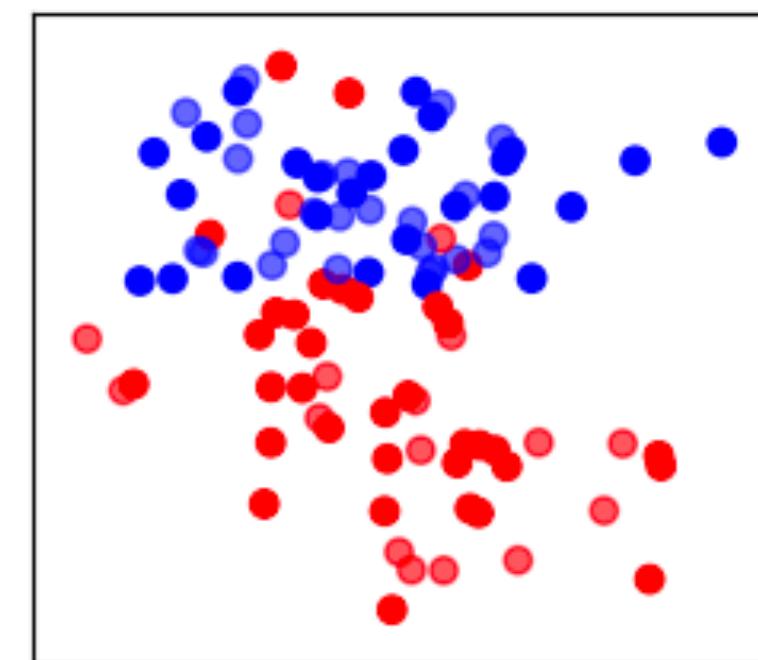
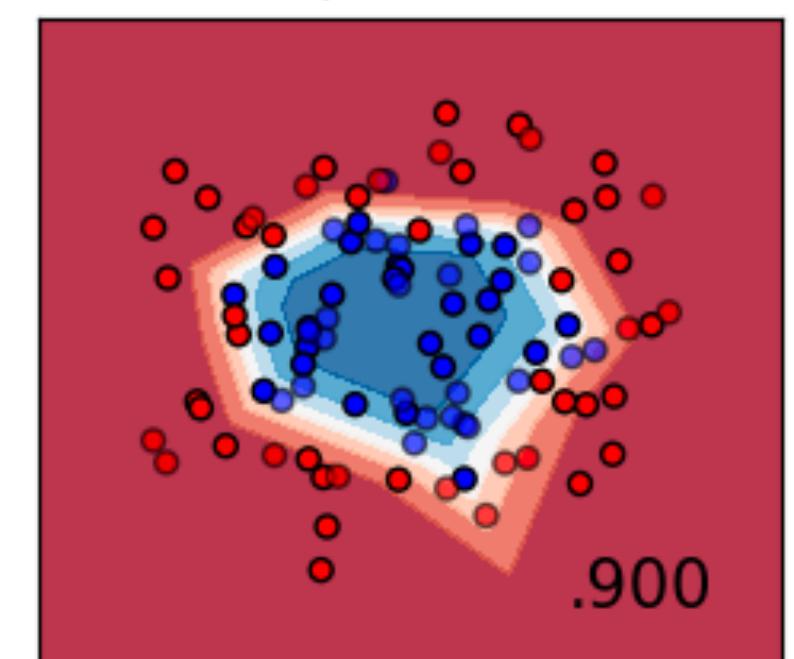
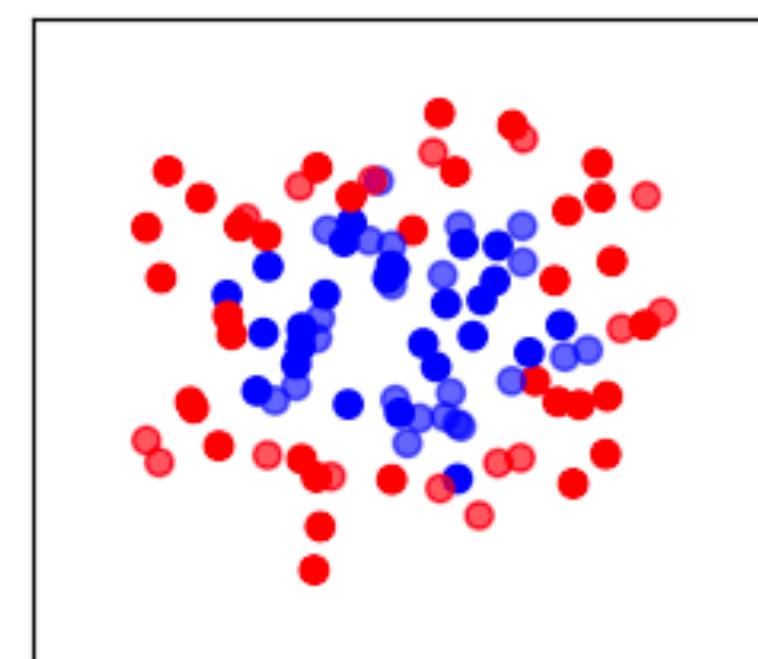
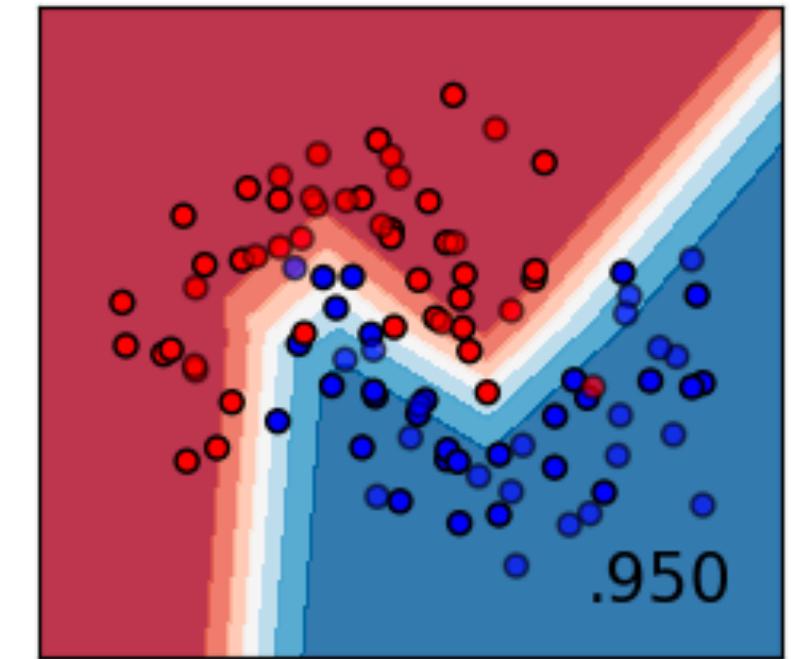
- Decision boundaries can correspond to logical rules, even when non-linear (e.g., XOR)
 - But most decision-boundaries are not easily explainable using symbolic operations
 - Rather, the feature space is carved up based on *similarity* to trained exemplars
- Decisions can be *probabilistic* rather than all-or-nothing category membership, based on the activation of the final layer



Training data

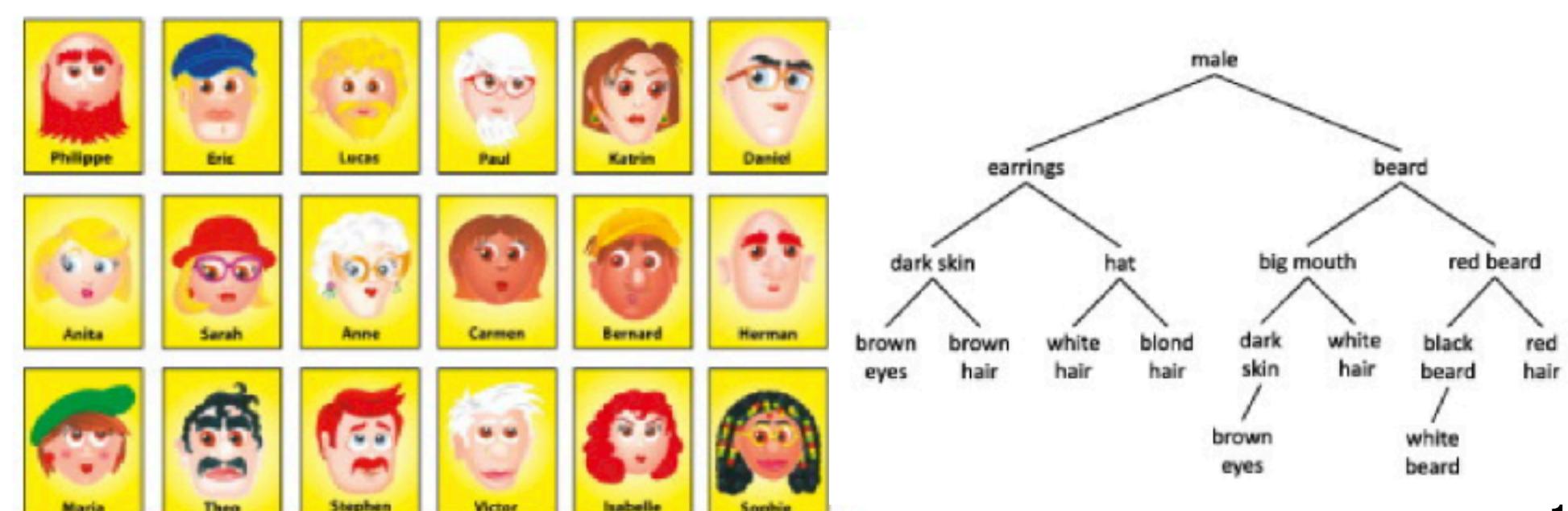
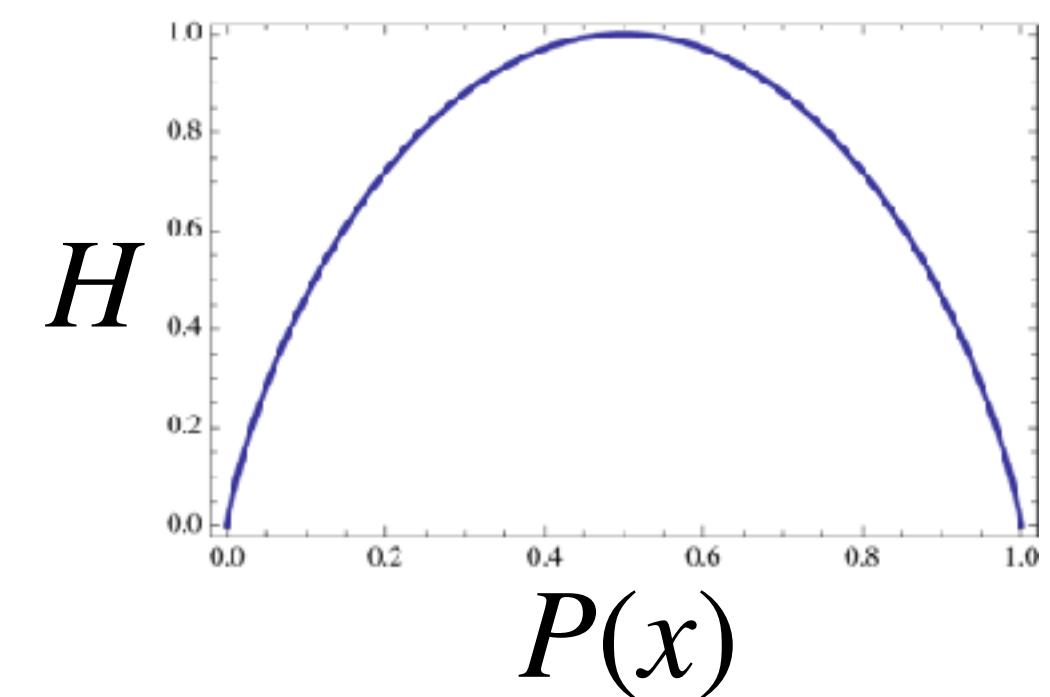
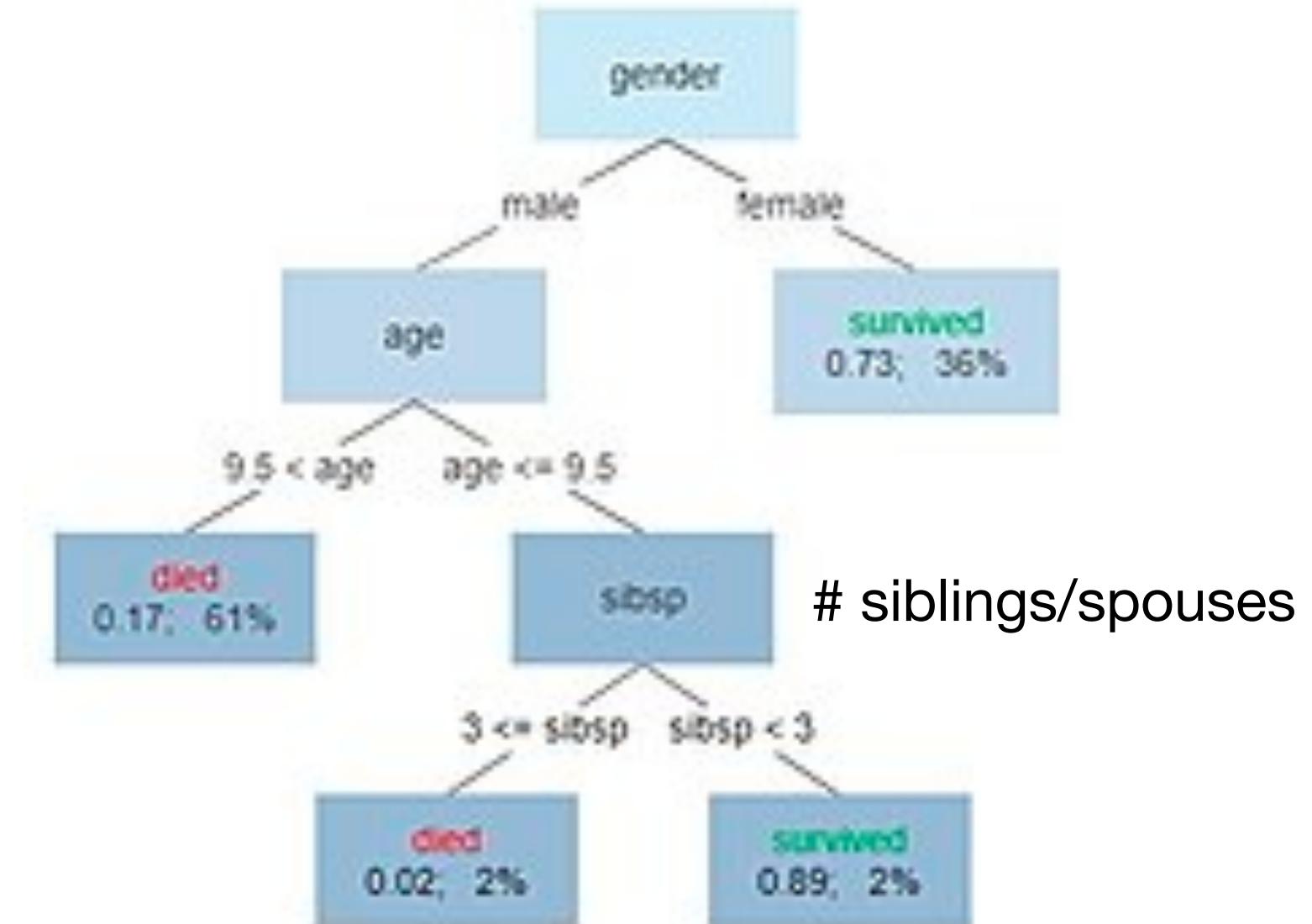


Decision boundary



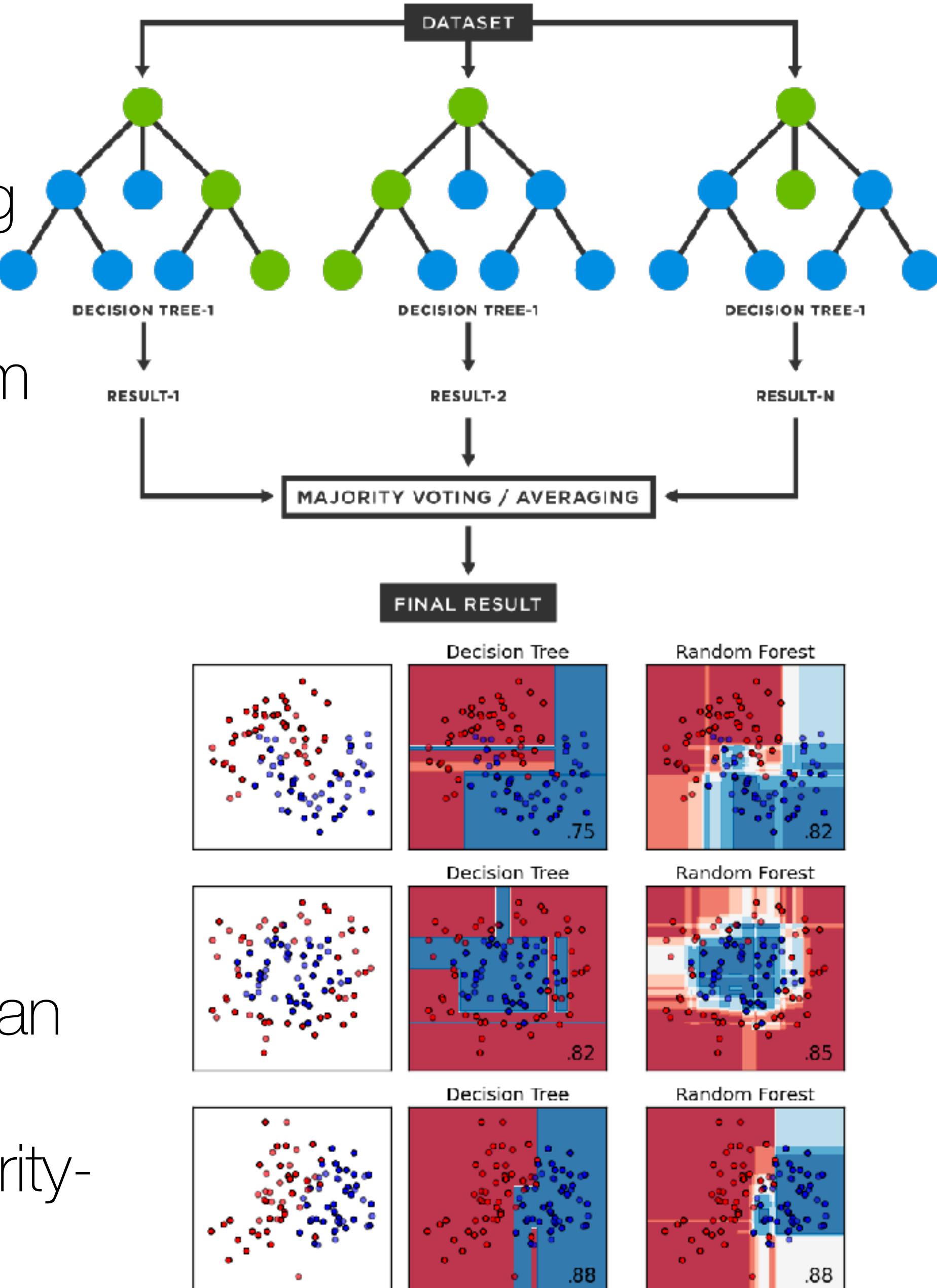
Decision-trees

- **Decision Trees** are the quintessential rule-based classifier
 - Easy to interpret, but can be prone to bias and overfitting
 - ID3 algorithm: Build the tree by maximizing *Information gain*
 - $IG(\mathbf{X}, f) = H(\mathbf{X}) - H(\mathbf{X} | f)$
 - where $H(\mathbf{X}) = - \sum P(x) \log P(x)$ is Shannon Entropy
 - How much does feature f reduce entropy?
 - The more the feature can even split the data (across labels), the greater the reduction of entropy and the greater the IG
 - If not naturally a binary feature, define a threshold that maximizes Entropy (e.g., 9.5 yrs of age)
 - Make a decision node using the feature with $\max(IG)$
 - Repeat until we run out of features



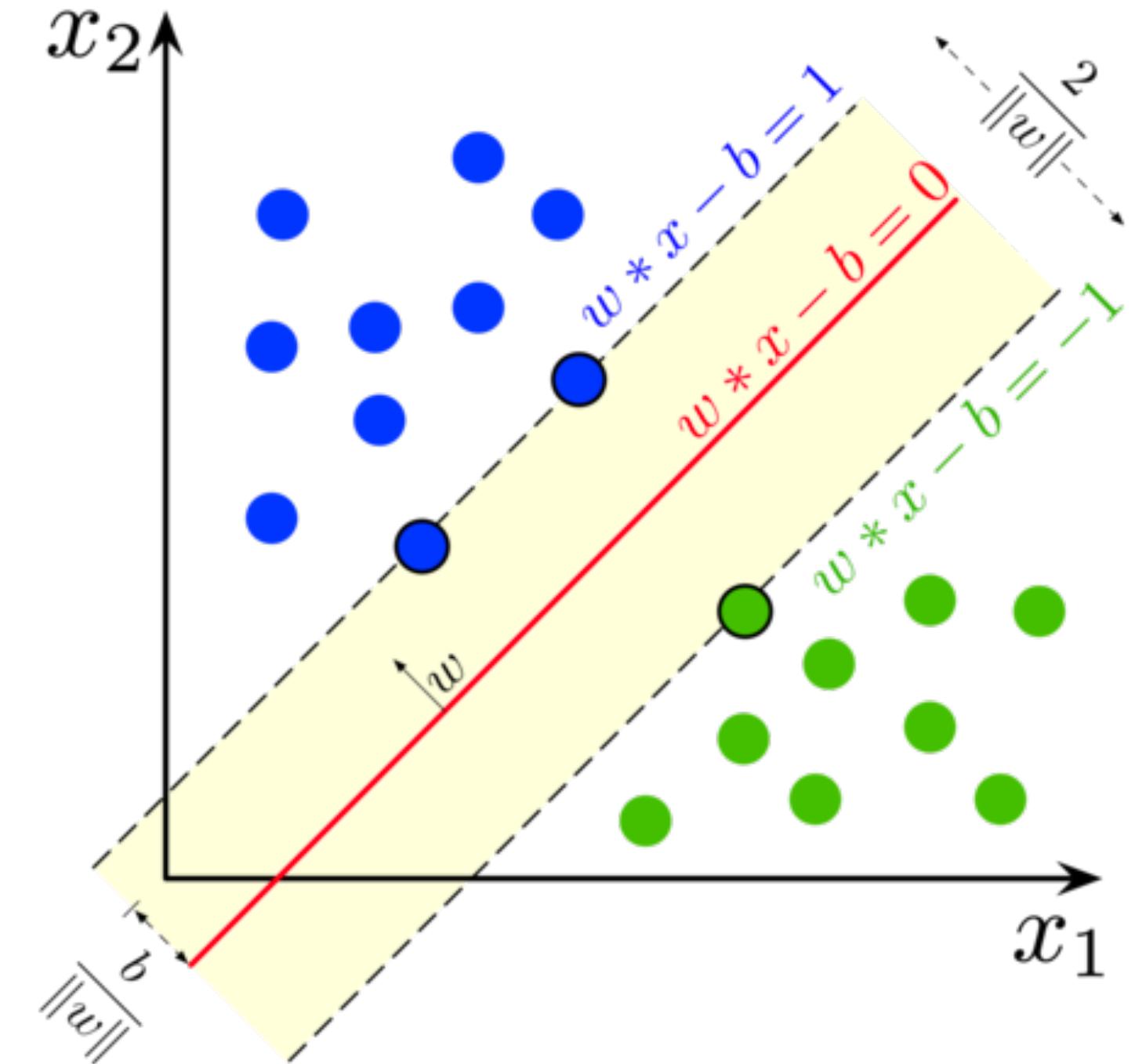
Random forests

- **Random forests** are an ensemble method combining random, uncorrelated decision trees
 - Each tree uses “feature bagging” to sample a random subset of features, ensuring low correlation among trees
 - Voting or averaging to make the final decision
- Ensemble methods are common in ML
 - Do brains also combine “opinions” from multiple decision-making systems?
- Aggregation over multiple trees is similar to how Bayesian concept learning operates over a distribution of rules, producing generalization patterns consistent with similarity-based theories



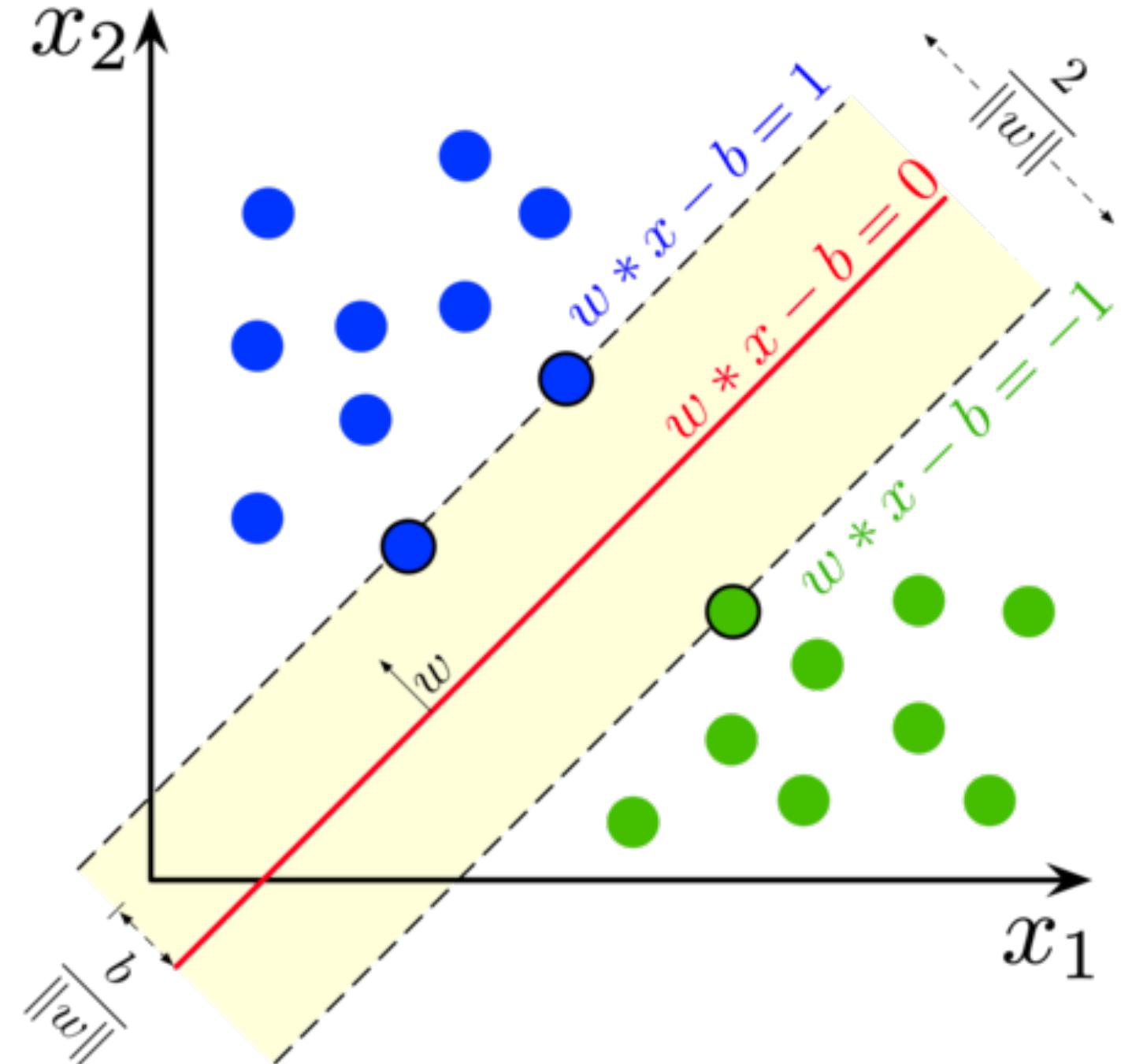
Support Vector Machines

- Learn a **decision boundary** $w^\top x - b = 0$ that best separates the data
- Let's start with the simplest setting, where we assume the data is linearly separable
- **Hard-margin**: with $y_i \in [-1, 1]$, we want
 - $y_i(w^\top x - b) \geq 1$ (i.e., all data classified correctly)
 - And to maximize the **margin** between classes $\frac{2}{\|w\|}$, which we do by minimizing $\|w\|$
 - This gives us a constrained optimization problem:
$$\mathcal{L}(w, b) = \|w\| \text{ subject to } y_i(w^\top x - b) \geq 1$$
 - The solution is completely determined by the x_i closest to the decision-boundary (i.e., support vectors)

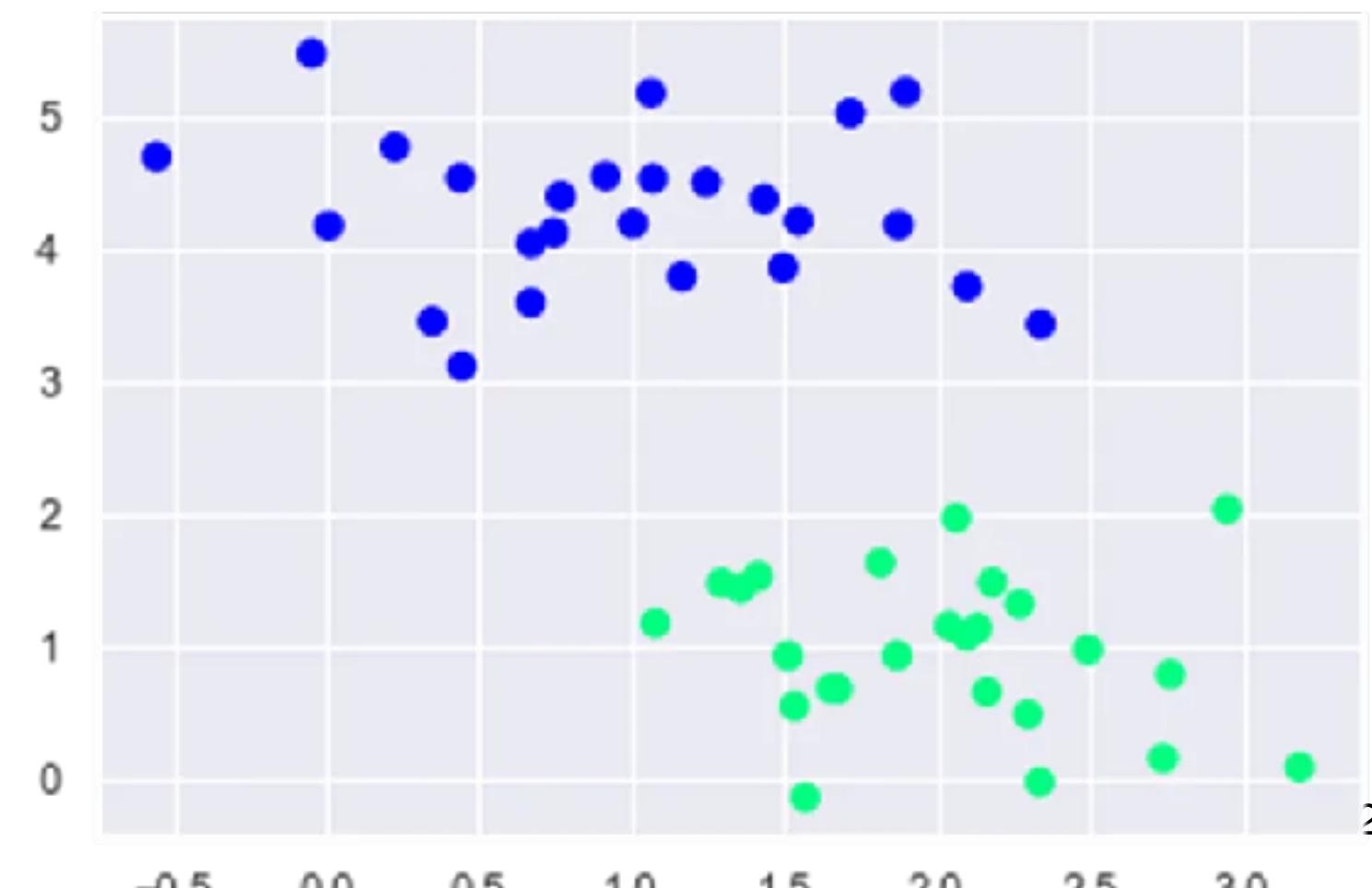


Support Vector Machines

- Learn a **decision boundary** $w^\top x - b = 0$ that best separates the data
- Let's start with the simplest setting, where we assume the data is linearly separable
- **Hard-margin**: with $y_i \in [-1, 1]$, we want
 - $y_i(w^\top x - b) \geq 1$ (i.e., all data classified correctly)
 - And to maximize the **margin** between classes $\frac{2}{\|w\|}$, which we do by minimizing $\|w\|$
 - This gives us a constrained optimization problem:
$$\mathcal{L}(w, b) = \|w\| \text{ subject to } y_i(w^\top x - b) \geq 1$$
 - The solution is completely determined by the x_i closest to the decision-boundary (i.e., support vectors)

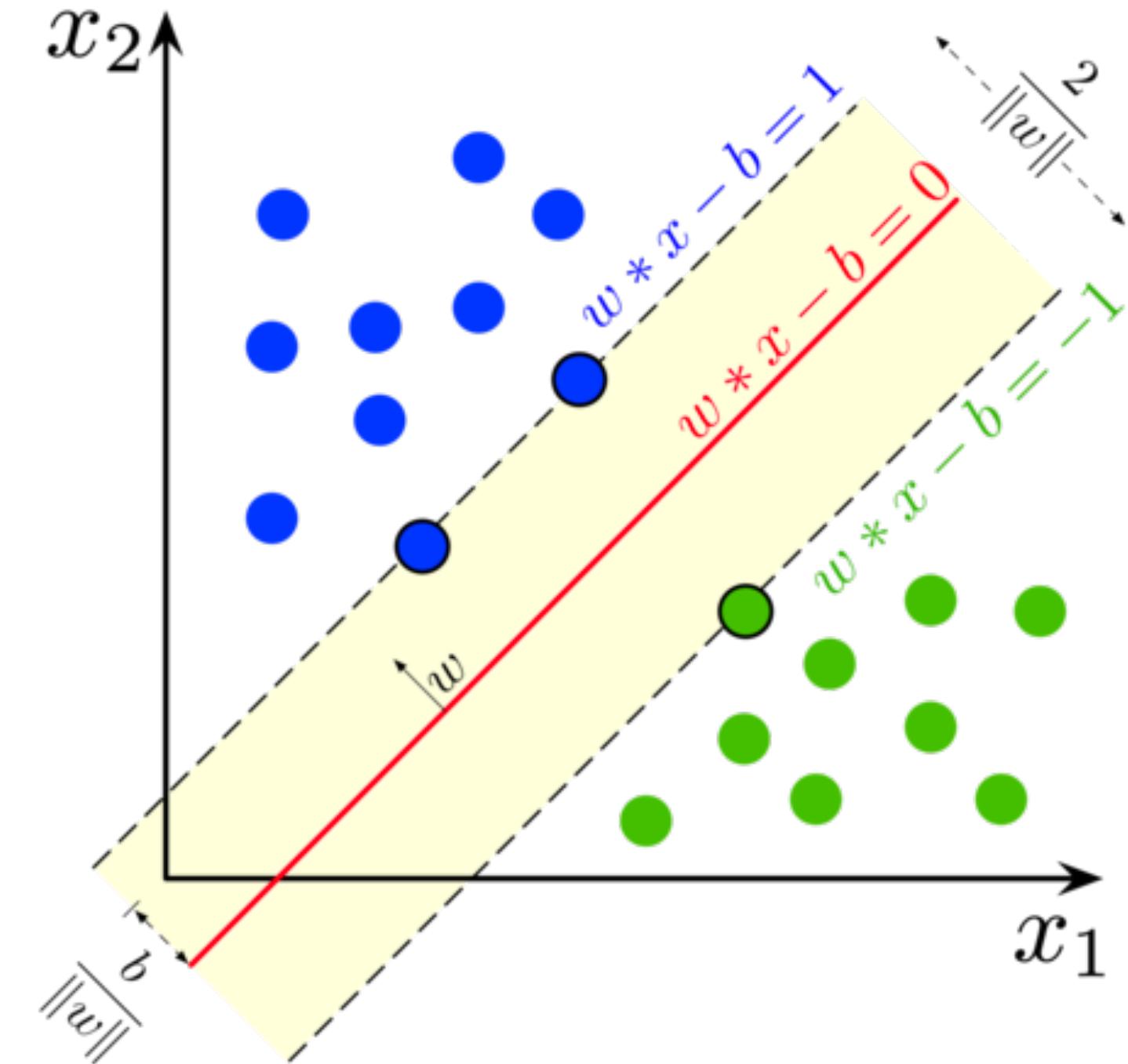


```
from sklearn.svm import SVC # "Support vector classifier"
model = SVC(kernel='linear', C=1) # By setting kernel= linear and C=1, we use hard margin
model.fit(X, y)
```

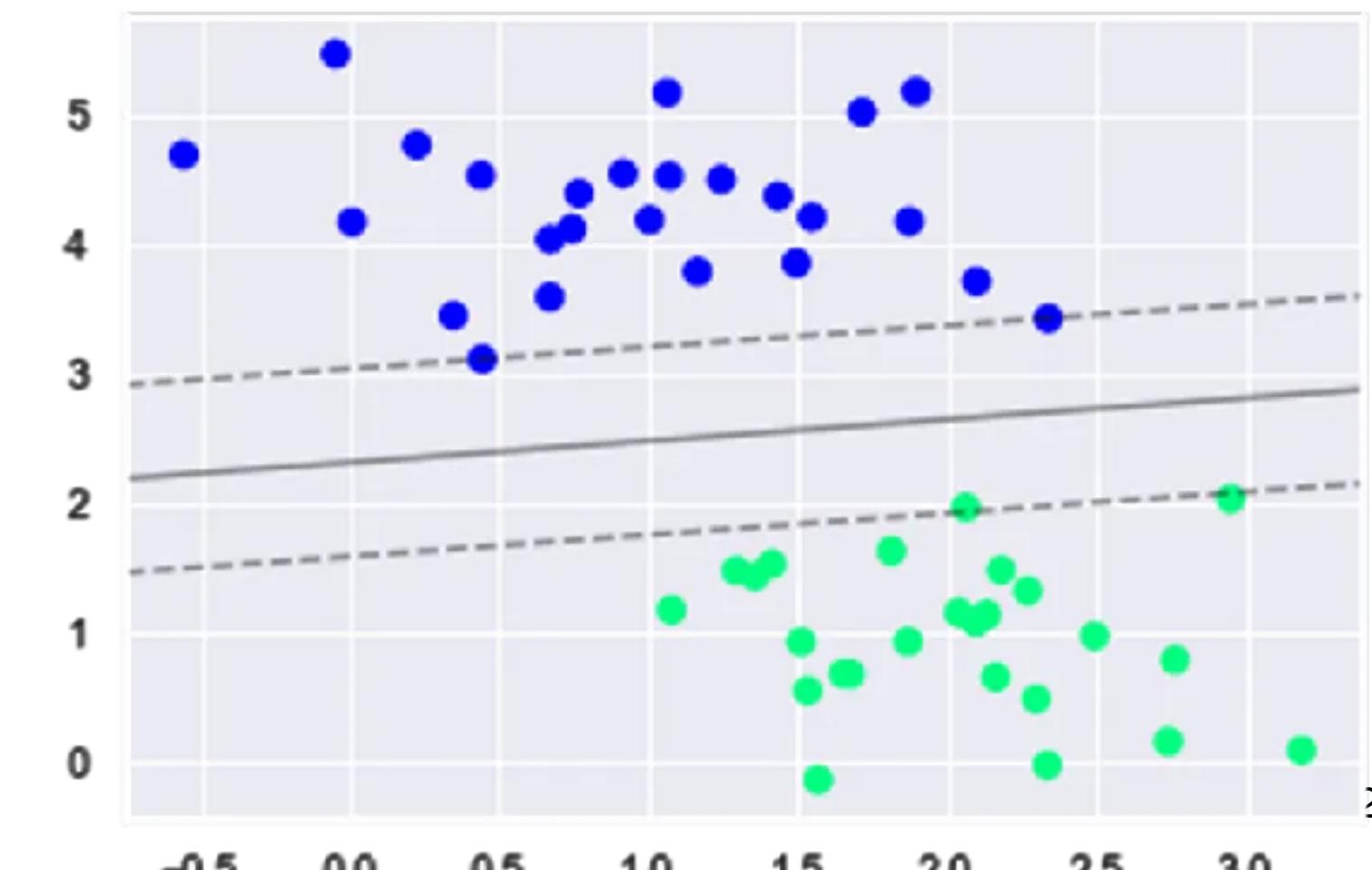


Support Vector Machines

- Learn a **decision boundary** $w^\top x - b = 0$ that best separates the data
- Let's start with the simplest setting, where we assume the data is linearly separable
- **Hard-margin**: with $y_i \in [-1, 1]$, we want
 - $y_i(w^\top x - b) \geq 1$ (i.e., all data classified correctly)
 - And to maximize the **margin** between classes $\frac{2}{\|w\|}$, which we do by minimizing $\|w\|$
 - This gives us a constrained optimization problem:
$$\mathcal{L}(w, b) = \|w\| \text{ subject to } y_i(w^\top x - b) \geq 1$$
 - The solution is completely determined by the x_i closest to the decision-boundary (i.e., support vectors)



```
from sklearn.svm import SVC # "Support vector classifier"
model = SVC(kernel='linear', C=1) # By setting kernel= linear and C=1, we use hard margin
model.fit(X, y)
```

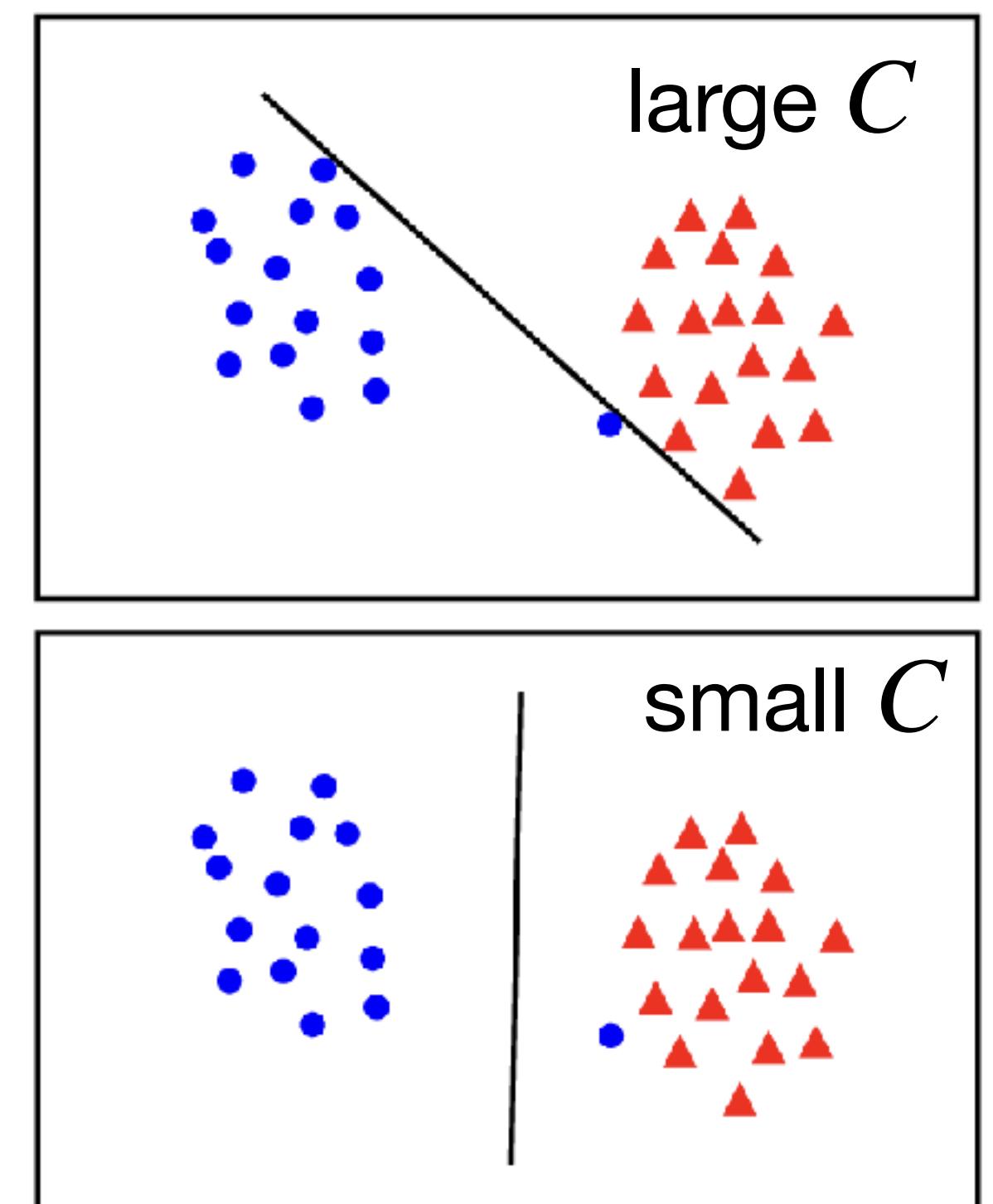
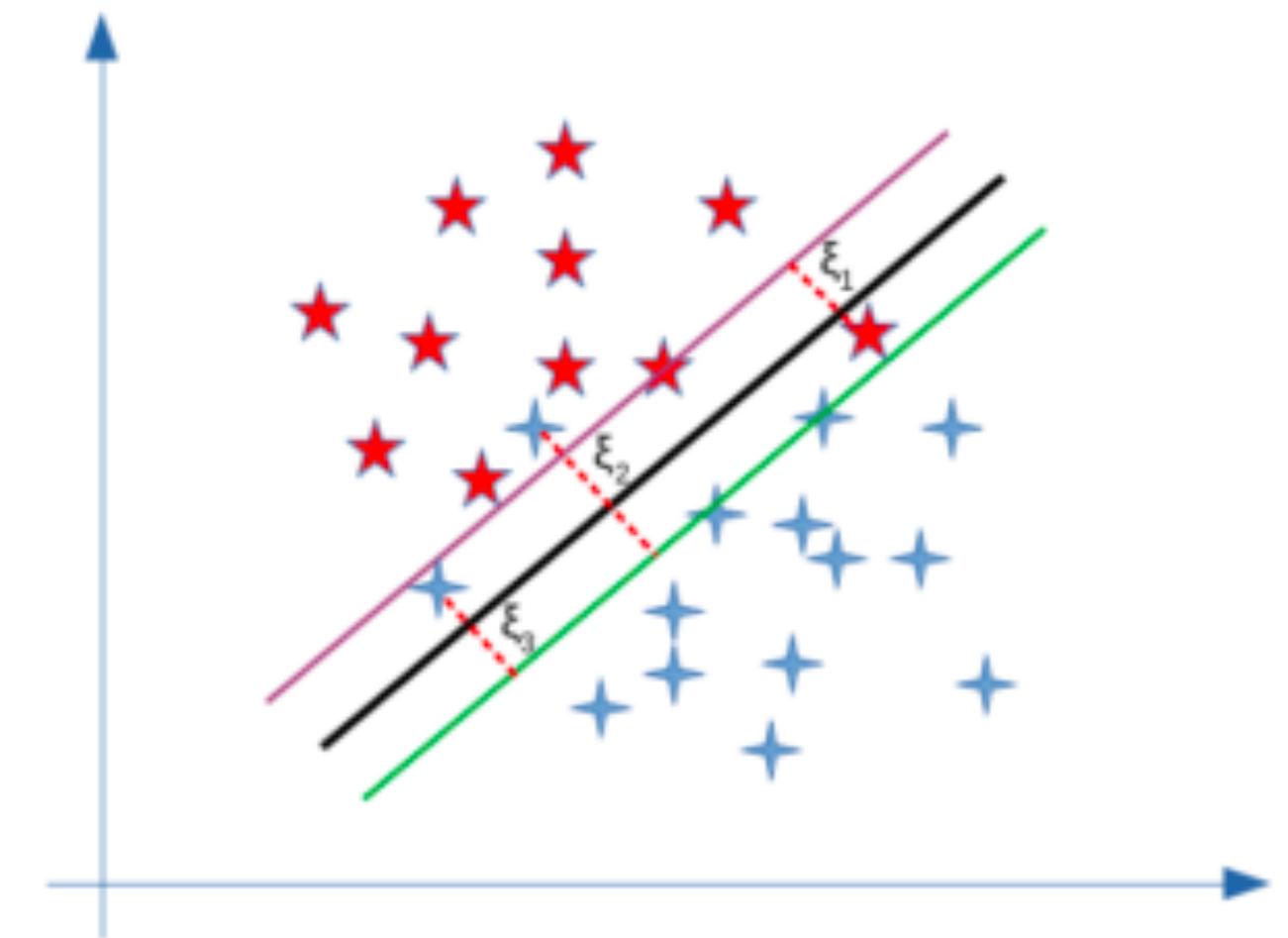


Support Vector Machines

- **Soft-margin:** Since data might not be linearly separable, use a soft-constraint

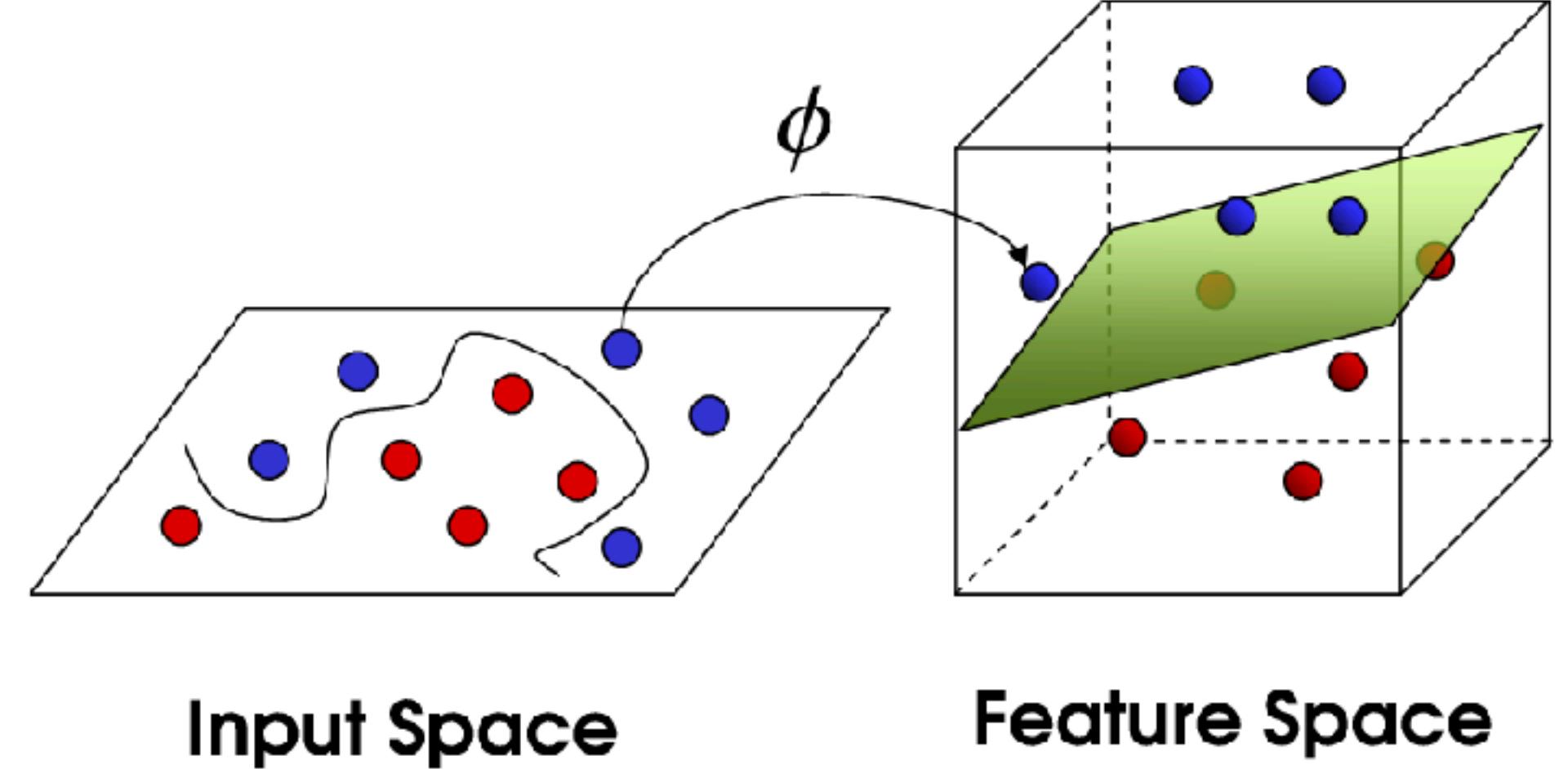
$$\mathcal{L}(w, b) = \|w\| - C \sum_i^N \zeta_i$$

- C is a penalty term defining how much we care about errors vs. a large margin
- Slack variable $\zeta_i = \max(0, 1 - y_i (w^\top x_i + b))$
 - $\zeta_i = 0$ for correctly classified points
 - $1 > \zeta_i > 0$ for margin violations
 - $\zeta > 1$ for incorrect classifications
- Smaller C allows for more errors in exchange for larger margins (and better generalization)



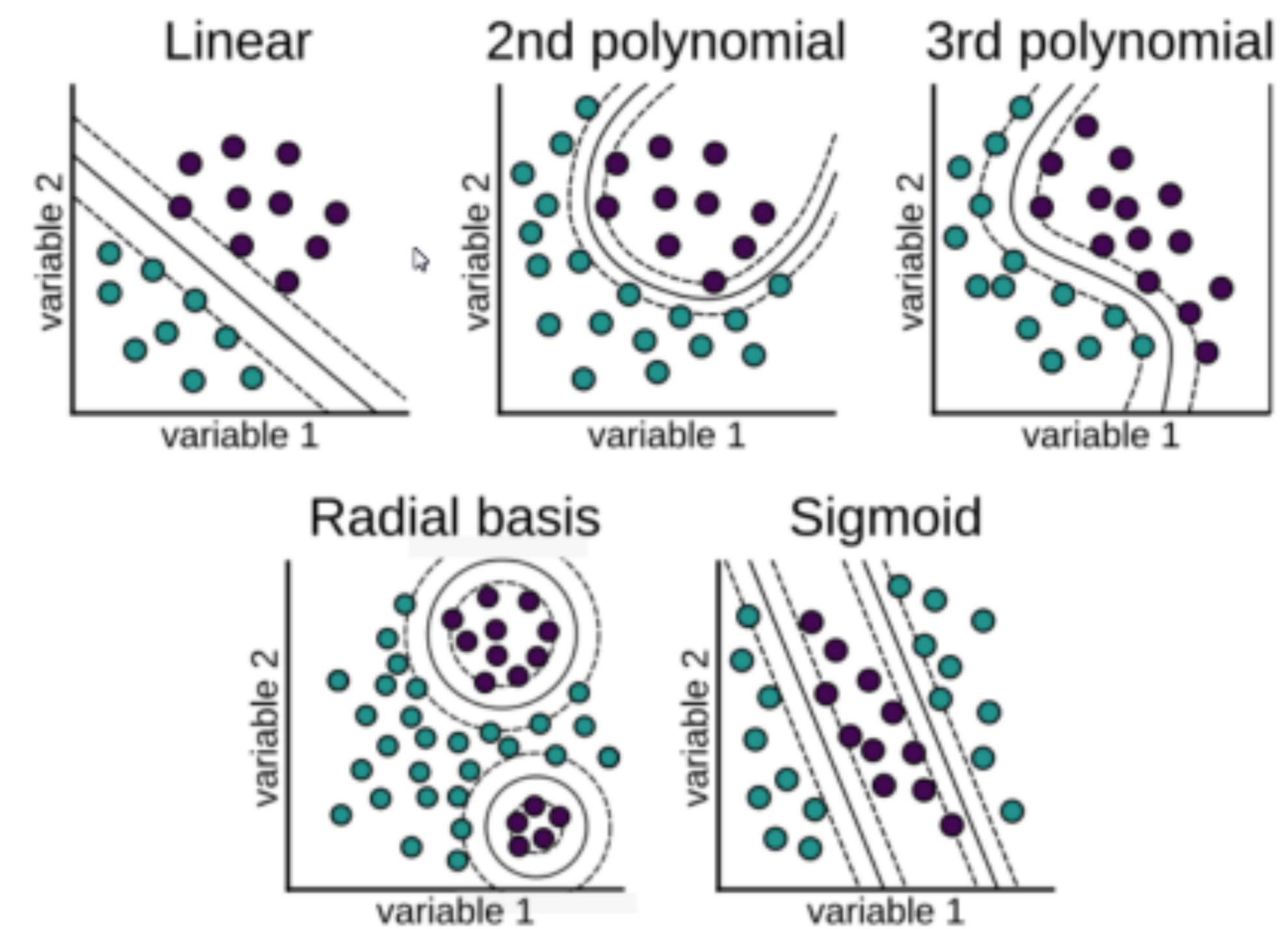
Kernel SVMs

- What about problems with non-linear decision boundaries?
- **Kernel trick** “projects” the data to a higher dimension, such that we can still learn a linear decision boundary
 - Rather than learning $w^T x - b = 0$, we use a kernel to map X onto a feature space $\Phi = \phi(X)$
e.g., polynomial kernel $\phi(x) = (1, x, x^2, x^3, \dots)$
 - We then substitute $\phi(x)$ for x and use all the same equations, e.g., decision boundary becomes
$$w^T \phi(x) - b = 0$$
 - There are many types of kernels, in fact, every neural network learned by gradient descent is approximately a kernel machine (i.e., $y = f(\phi(x))$); Domingos, 2020

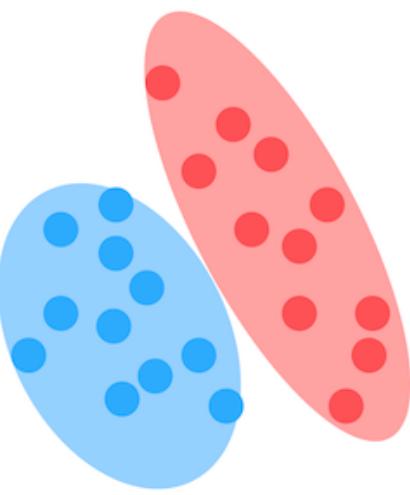


Input Space

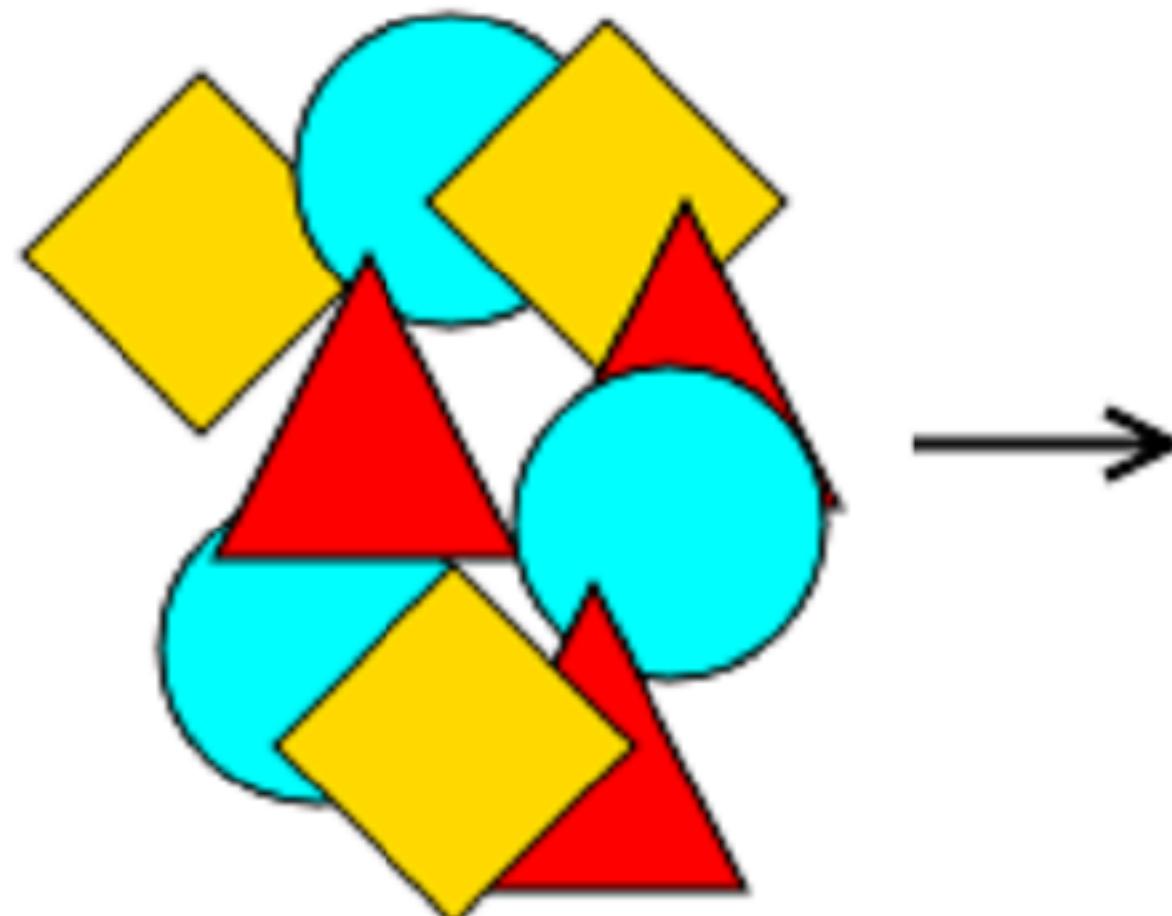
Feature Space



Naïve Bayes classifier



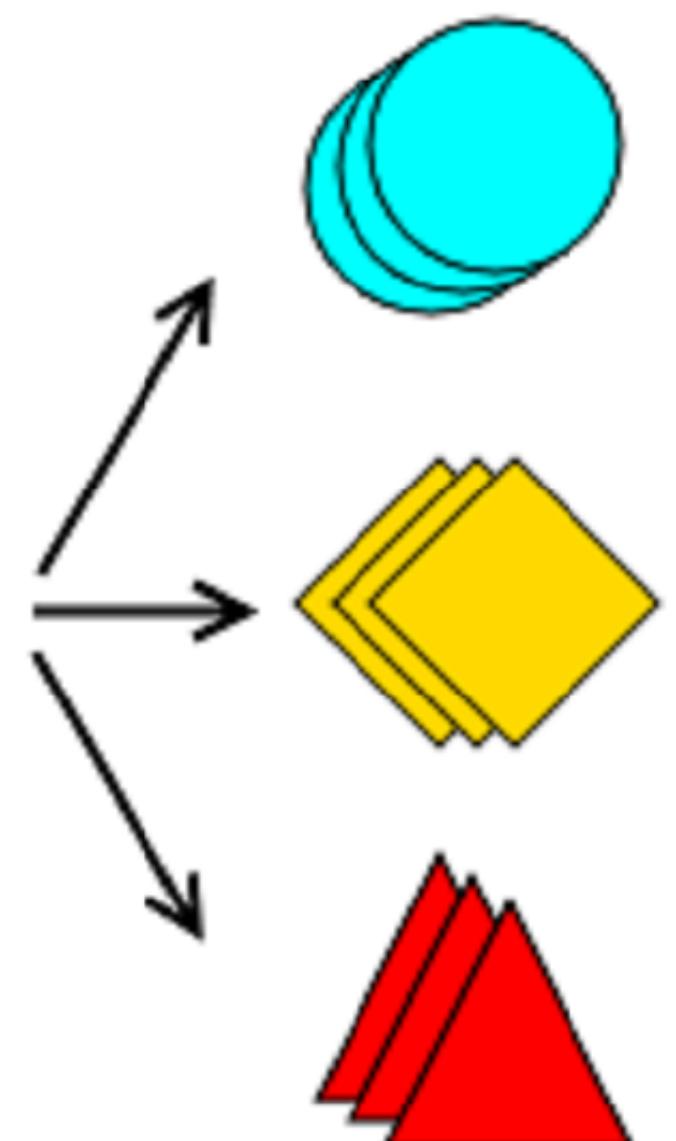
- First generative model: rather than learning a decision boundary, learns the distribution of the each category (which can be used to generate new data)
- Called naïve because we assume all features are independent
 - Easy and fast to learn
 - Can generalize to new feature values outside the data, although naïve assumption may be unrealistic
- We use Bayes' theorem to compute the posterior probability of an datapoint belonging to some class c_k given it's features \mathbf{x} :



$$P(c_k | \mathbf{x}) = \frac{P(\mathbf{x} | c_k)P(c_k)}{P(\mathbf{x})} \propto P(c_k) \prod_j^n P(x_j | c_k)$$

posterior = $\frac{\text{likelihood * class prior}}{\text{evidence}}$

denominator removed, because
it is the same for all data



$$P(c_k | \mathbf{x}) \propto P(c_k) \prod_j^n P(x_j | c_k)$$

Naïve Bayes classifier

- Computing the prior and likelihood
- **Prior** is just how frequent the category is in the data

$$P(c_k) = \text{count}(c_k)/N$$

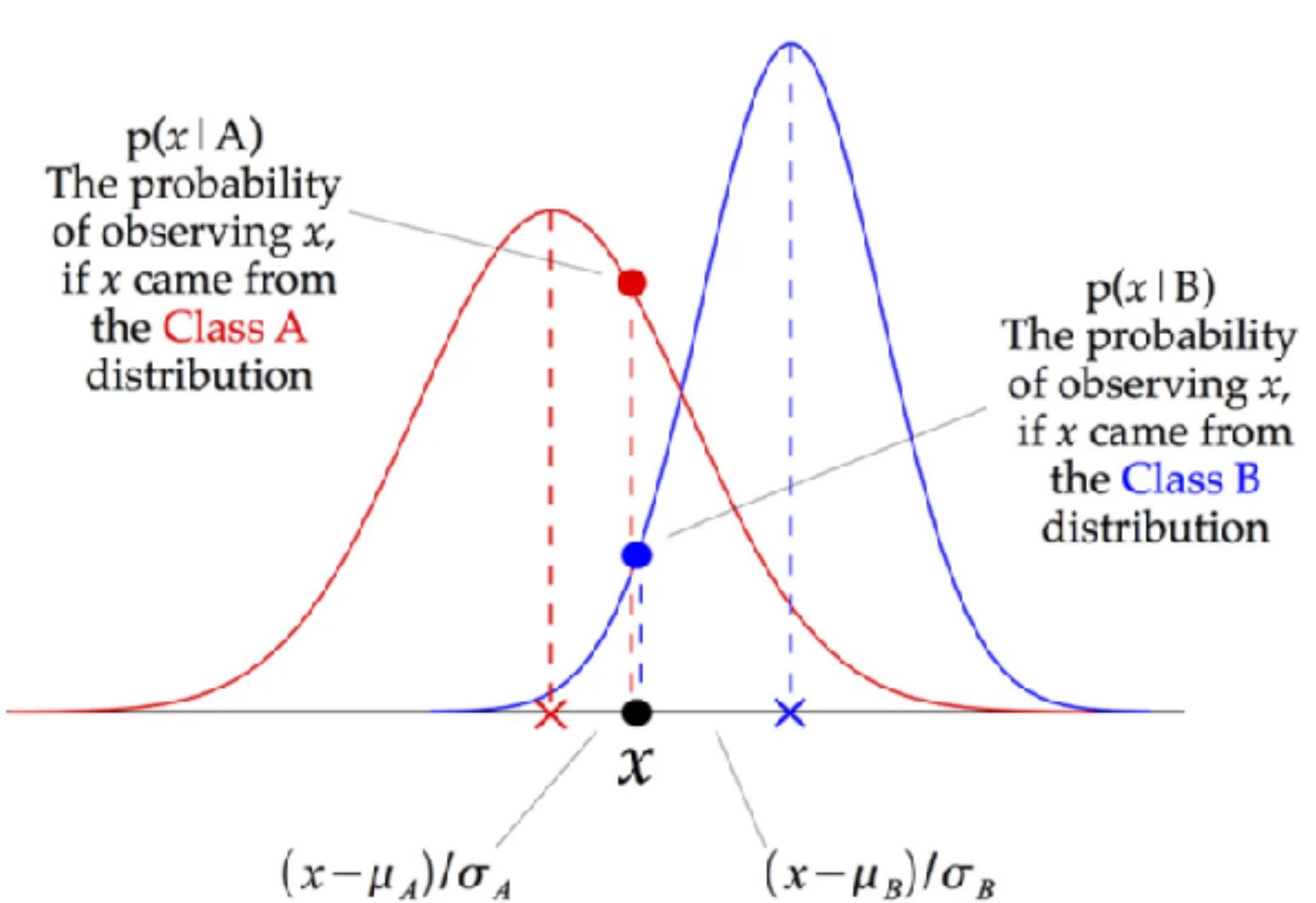
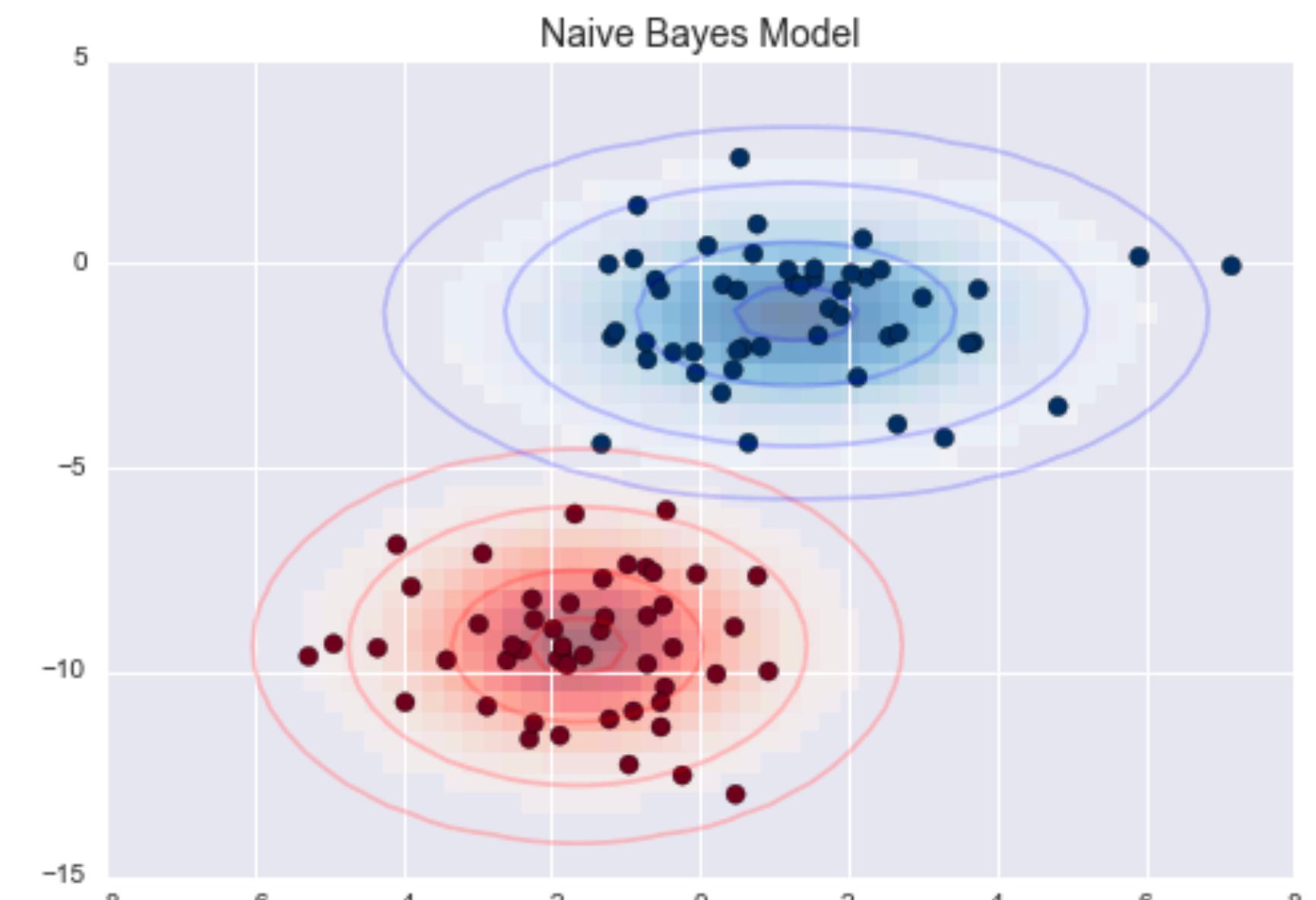
- **Likelihood:**

- When the **data is continuous**, we can assume a Gaussian distribution

$$P(\mathbf{x}_j | c_k) = \frac{1}{\sqrt{(2\pi)^d |\Sigma_k|}} \exp \left(-\frac{1}{2} (\mathbf{x}_j - \mu_k)^\top \Sigma_k^{-1} (\mathbf{x}_j - \mu_k) \right)$$

where μ_k and Σ_k are the mean vector and covariance matrix of the k-th category and d is the dimensionality of the data

- Likelihood is a function of distance from the mean of the c_k Gaussian



$$P(c_k | \mathbf{x}) \propto P(c_k) \prod_j^n P(x_j | c_k)$$

Naïve Bayes classifier

- For **text classification** (e.g., spam filters), where we have discrete variables, we can use a “bag of words” representation
- From data, extract vocabulary V , where each $\mathbf{x}_i = (x_1, x_2, \dots, x_V)$ represents the counts for each possible word
- Likelihood with Laplacian smoothing (to avoid divide by 0)

$$P(x_j | c_k) = \frac{\text{count}(x_j, c_k) + 1}{\sum_{x \in V} \text{count}(x, c_k) + 1}$$

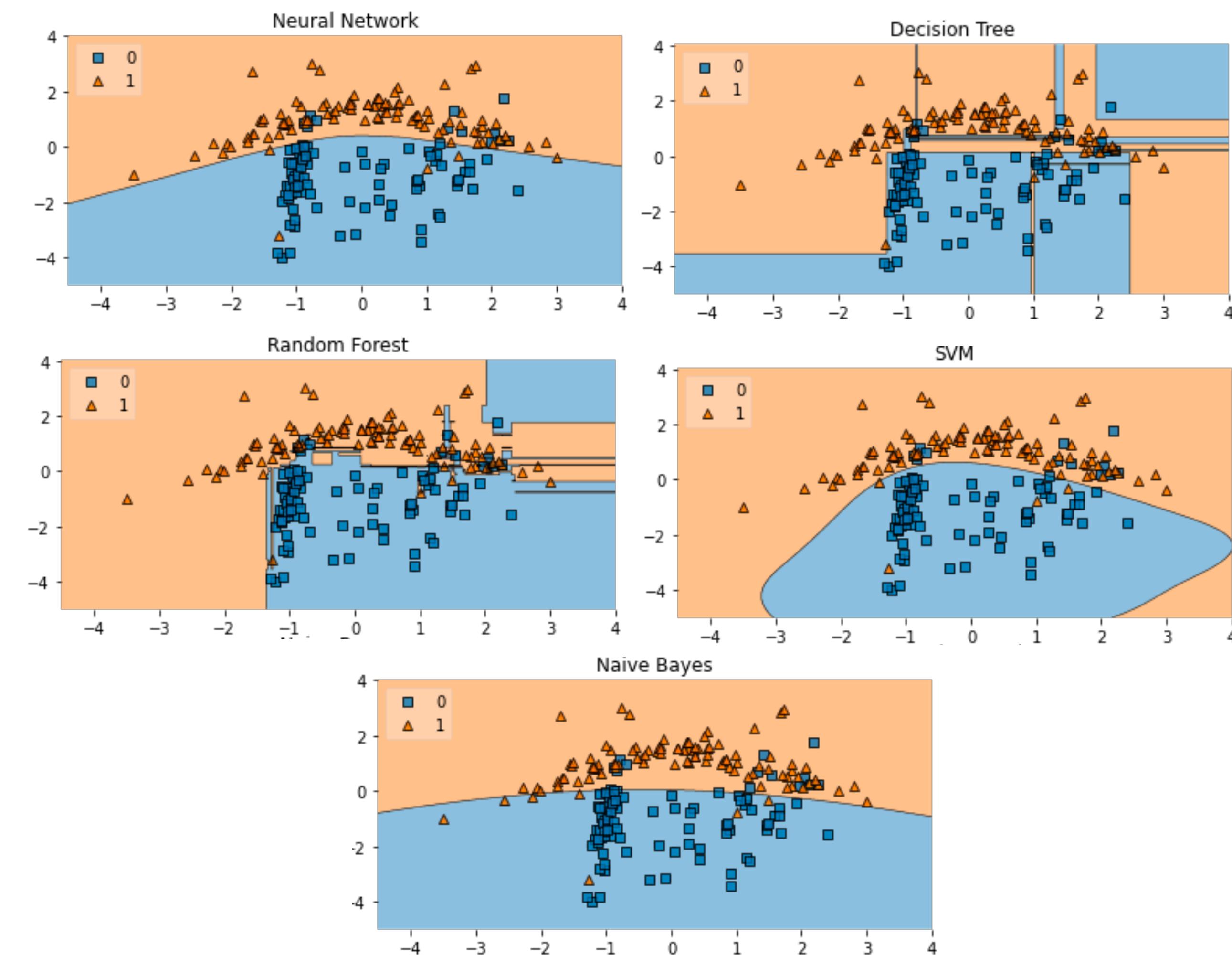


Frequency of word j within category

Frequency of all words

Supervised learning summary

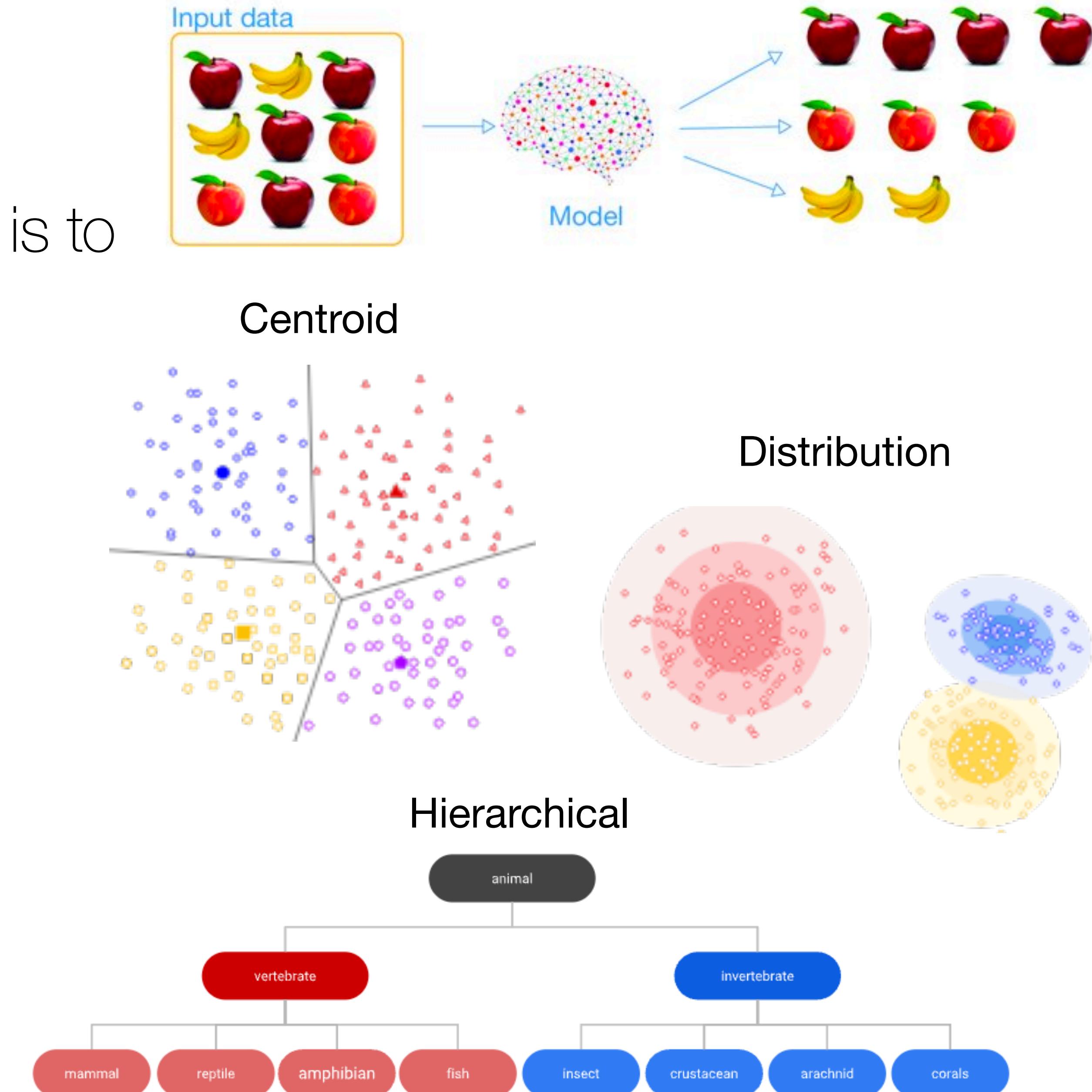
- Supervised learning is a **classification** problem
- Each method yields a corresponding **decision boundary** (rule-based interpretation)
- However, only decision trees operate on explicit rules
- Most **discriminative** approaches use similarity-based mechanisms (e.g., MLPs and SVMs) to arrive at a decision-boundaries based on carving up self-similar regions based on labeled exemplars
- However, **generative** methods learn explicit representations of each category
 - Naïve Bayes learns distributions for each category (prototype interpretation)



5 min break

Unsupervised learning

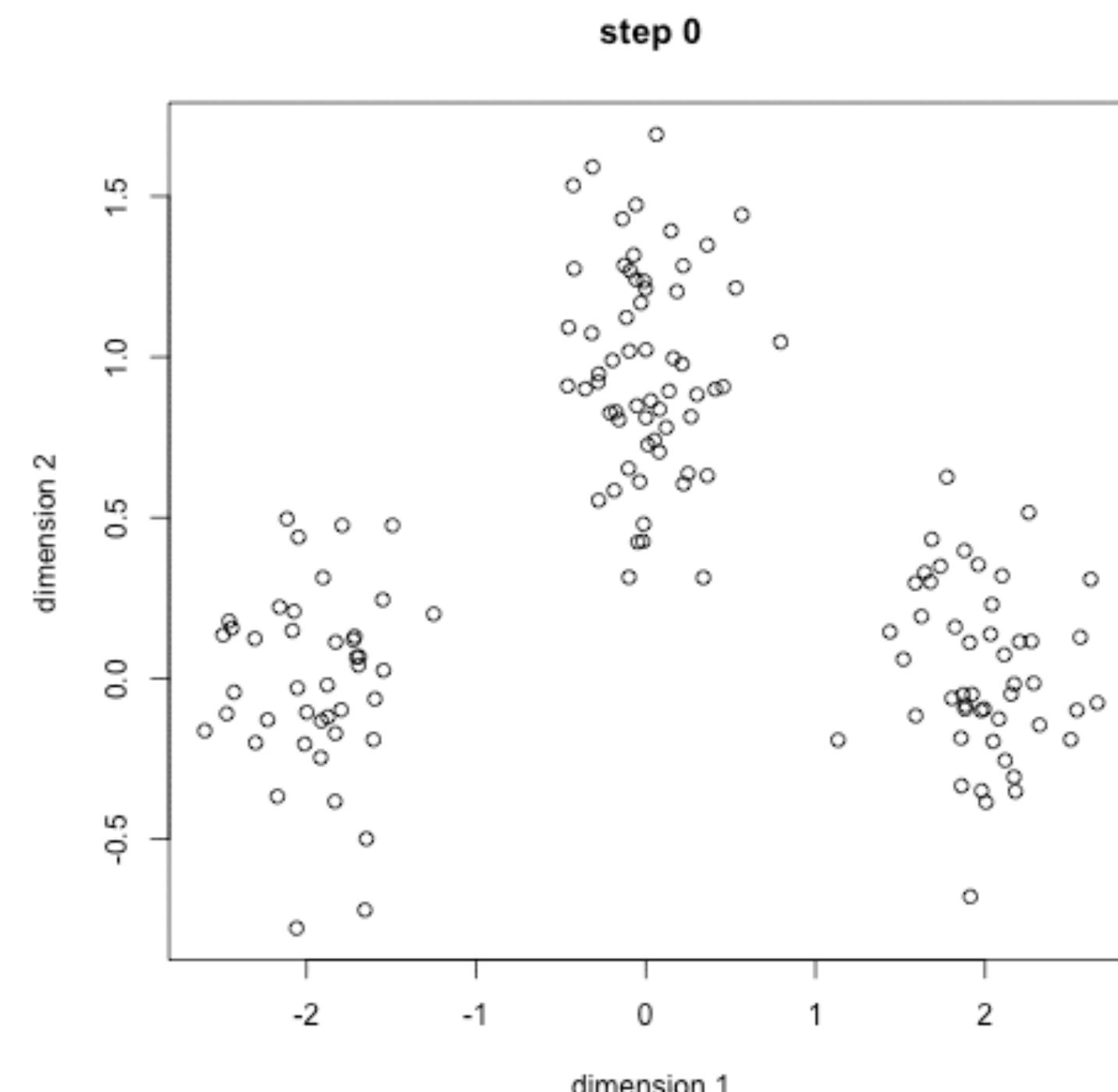
- Without supervised labels, the goal is to learn clusters based on similarity
- Types of clustering algorithms:
 - Centroid-based clustering
 - e.g., k-means
 - Distribution-based clustering
 - e.g., Gaussian mixture models
 - Hierarchical clustering
 - e.g., Agglomerative



k -means clustering

Learn k centroids that minimize within-cluster variance

1. Pick the number of clusters k
2. Randomly select the centroid for each cluster
3. Assign all points to the closest centroid
4. Recompute centroid based on assigned points (i.e., mean)
5. Repeat until centroids do not change or max number of iterations reached

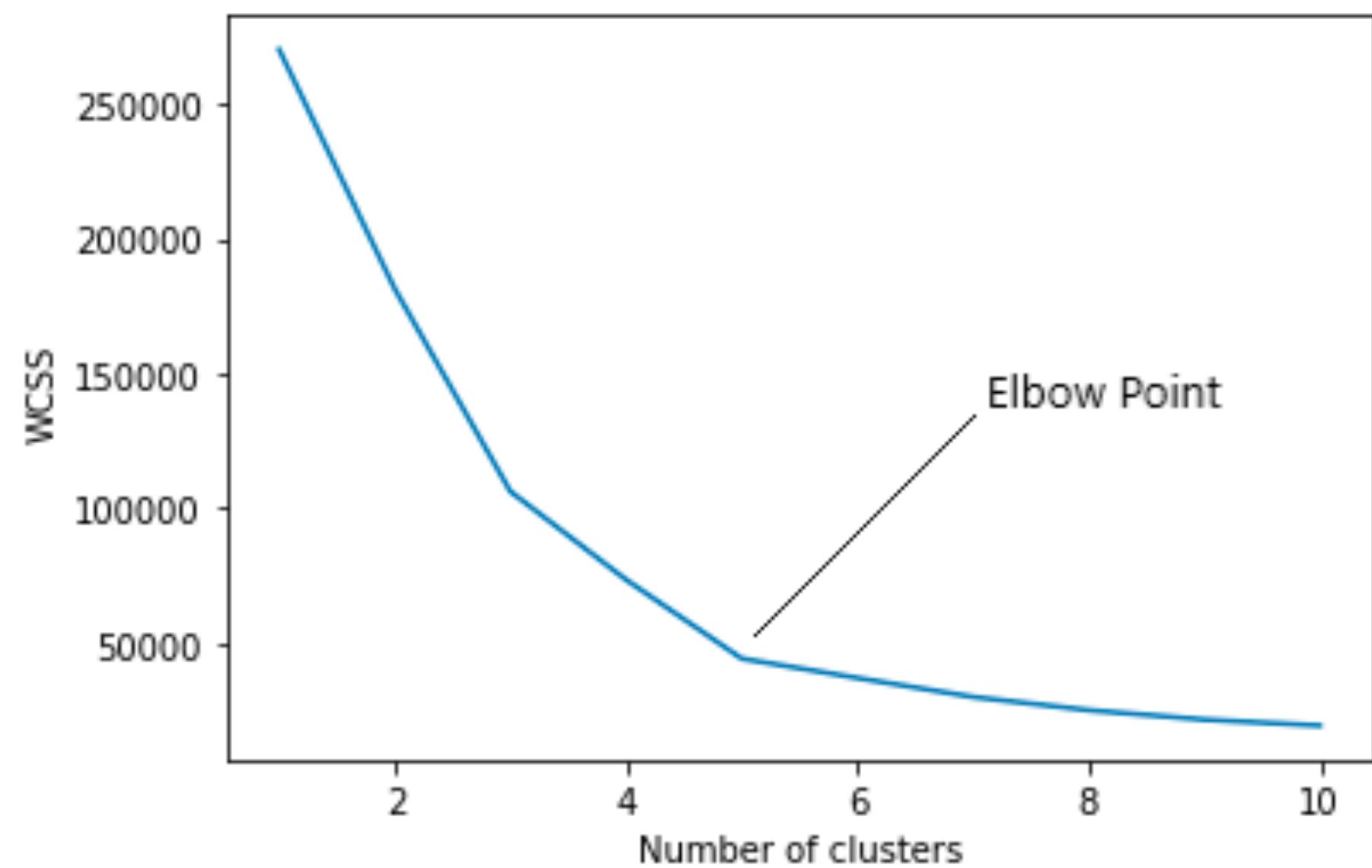


How do we pick the number of clusters?

Elbow method:

- Within-cluster sum of squares (WCSS) as a goodness-of-fit metric:
for each cluster $c \in 1, \dots, k$ compute the squared distance from each assigned datapoint x_i to the centroid μ_c

$$WCSS = \sum_{c=1}^k \sum_{i=1}^m (x_i - \mu_c)^2$$

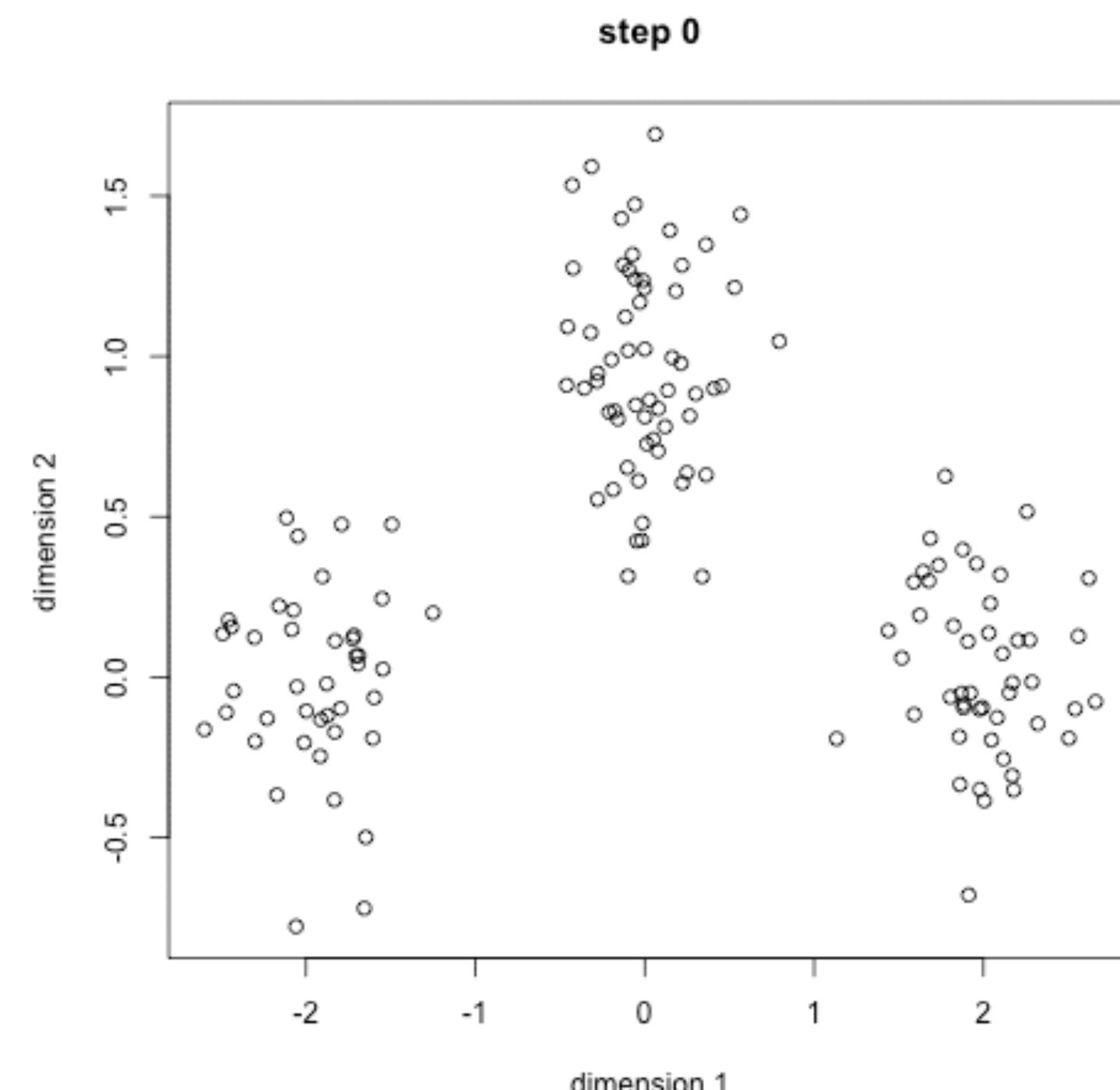


- Pick the number of clusters where WCSS begins to level off

k -means clustering

Learn k centroids that minimize within-cluster variance

1. Pick the number of clusters k
2. Randomly select the centroid for each cluster
3. Assign all points to the closest centroid
4. Recompute centroid based on assigned points (i.e., mean)
5. Repeat until centroids do not change or max number of iterations reached

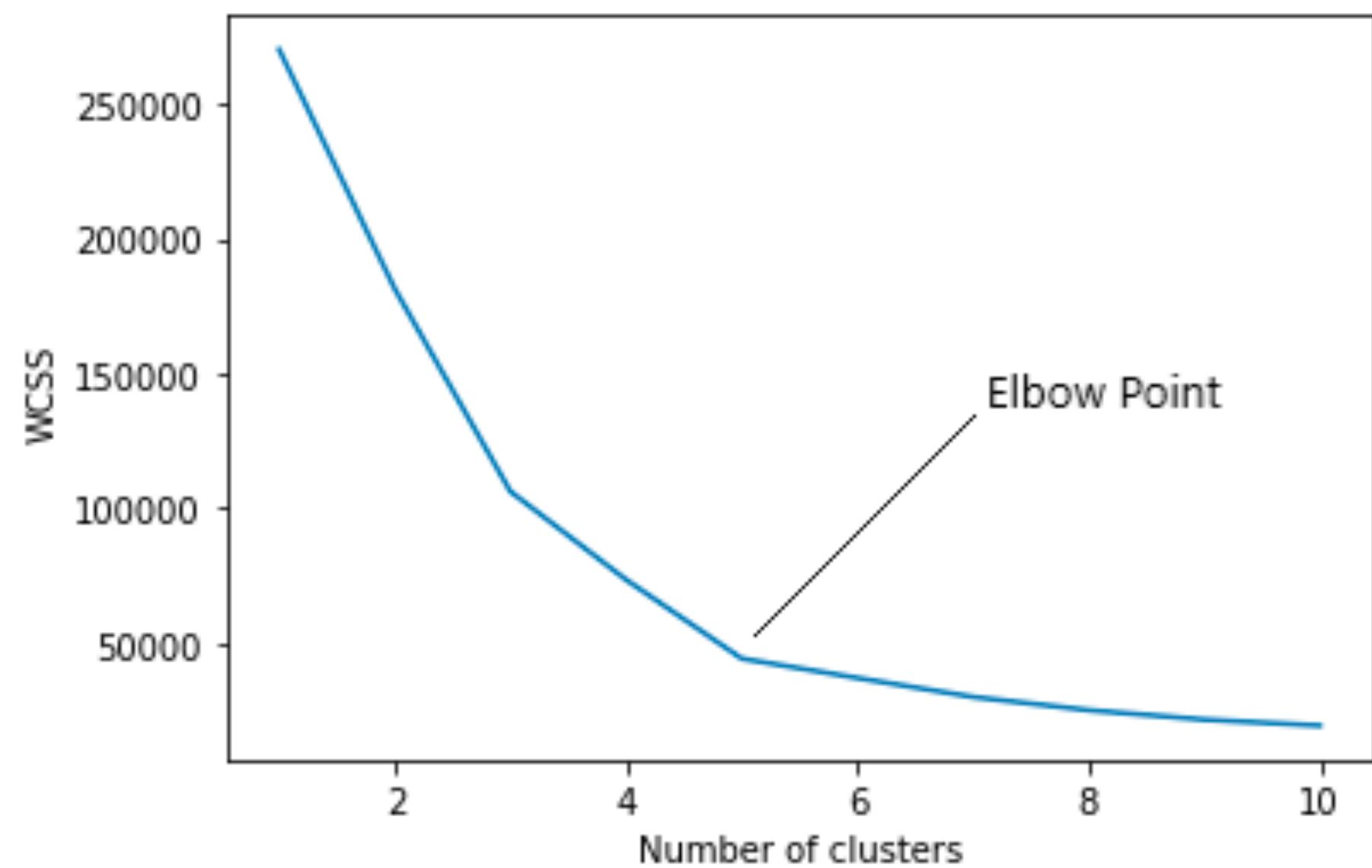


How do we pick the number of clusters?

Elbow method:

- Within-cluster sum of squares (WCSS) as a goodness-of-fit metric:
for each cluster $c \in 1, \dots, k$ compute the squared distance from each assigned datapoint x_i to the centroid μ_c

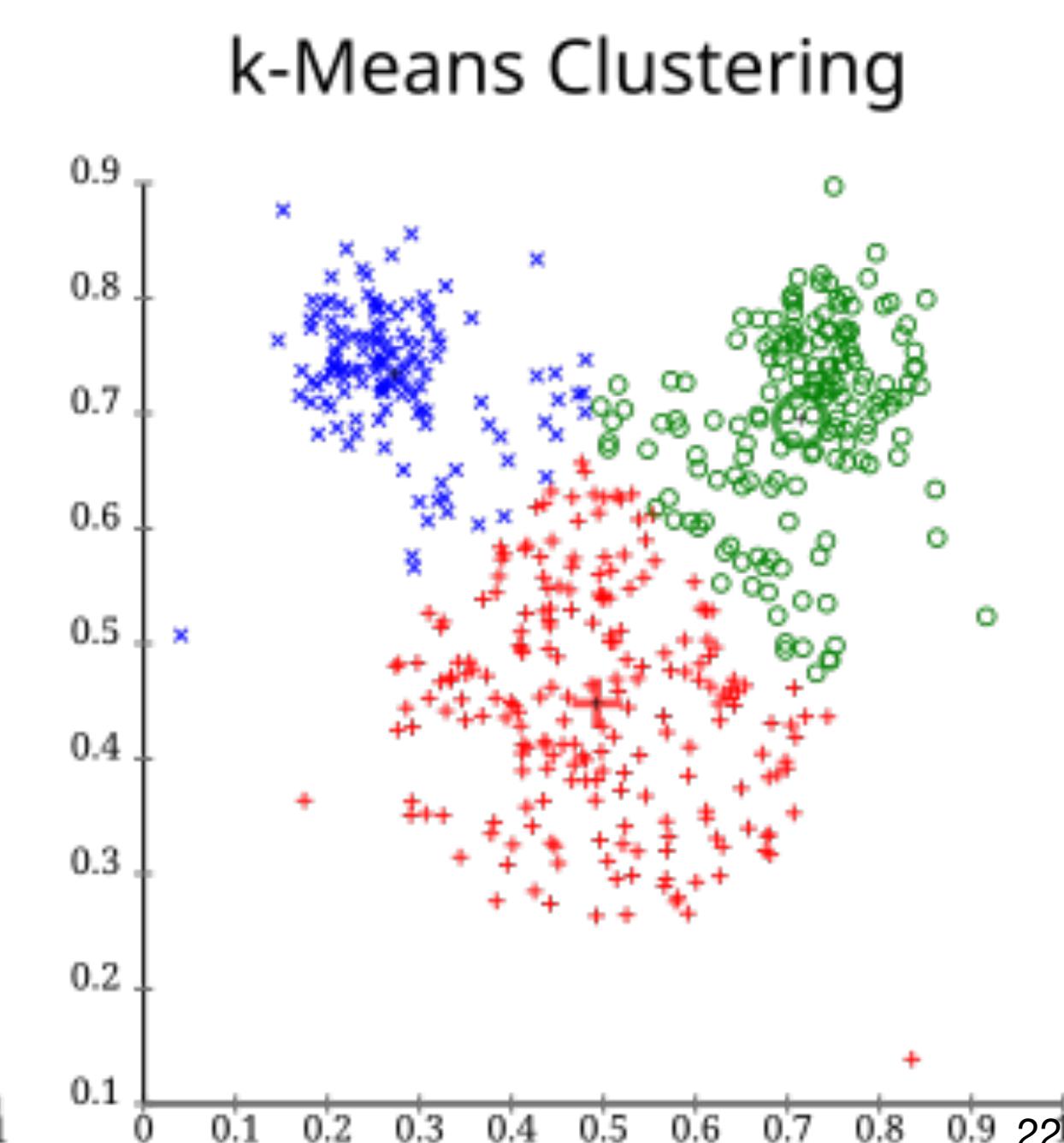
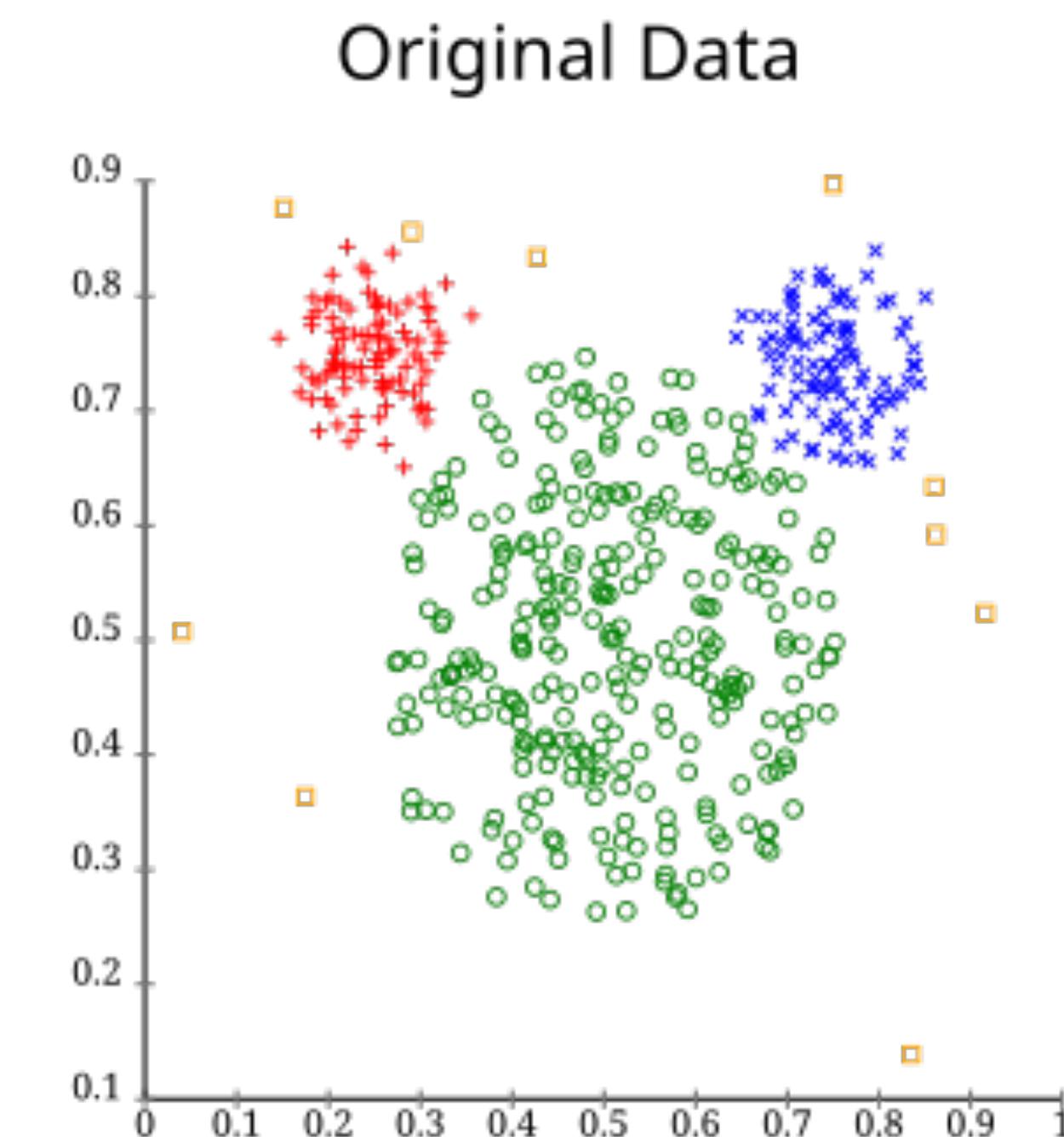
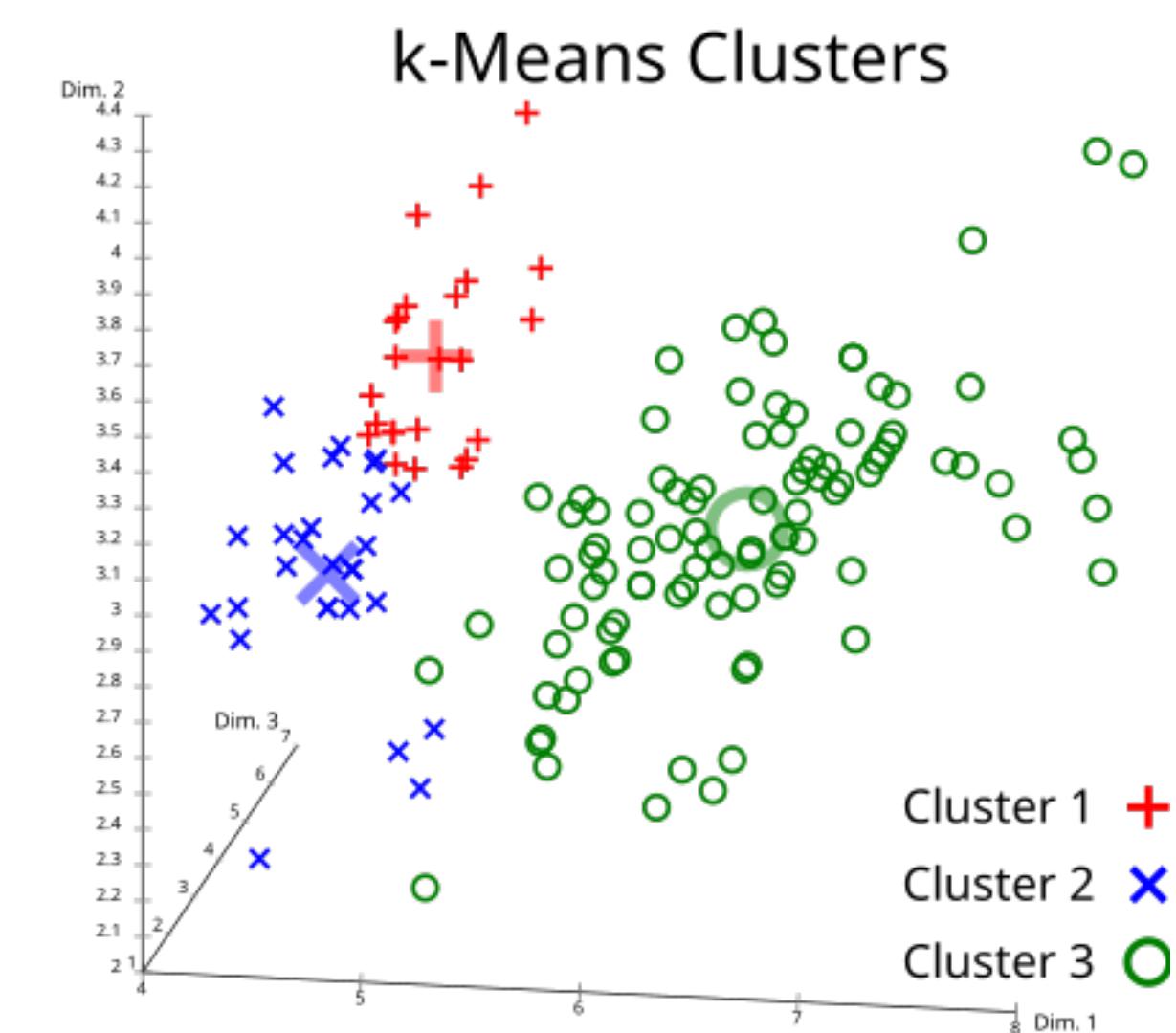
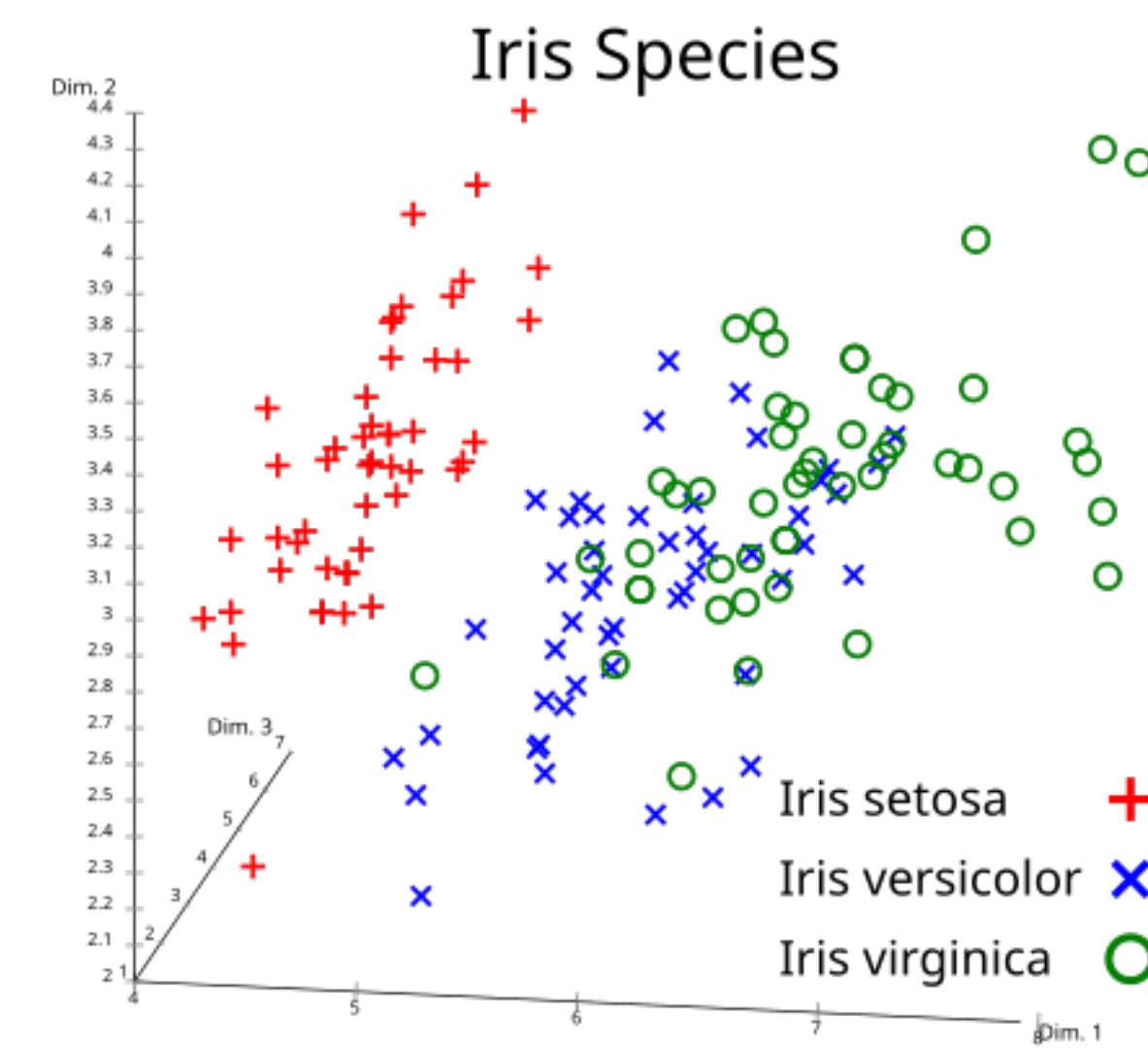
$$WCSS = \sum_{c=1}^k \sum_{i=1}^m (x_i - \mu_c)^2$$



- Pick the number of clusters where WCSS begins to level off

k-means clustering

- Main limitations are also what makes it efficient
 - Assumption of spherical clusters
 - The centroid is defined by the Euclidean mean
 - Assumption that clusters are of similar size
 - Assignment of data to the nearest cluster
 - These simplifying assumptions mean that that it works well on some datasets, but not on others

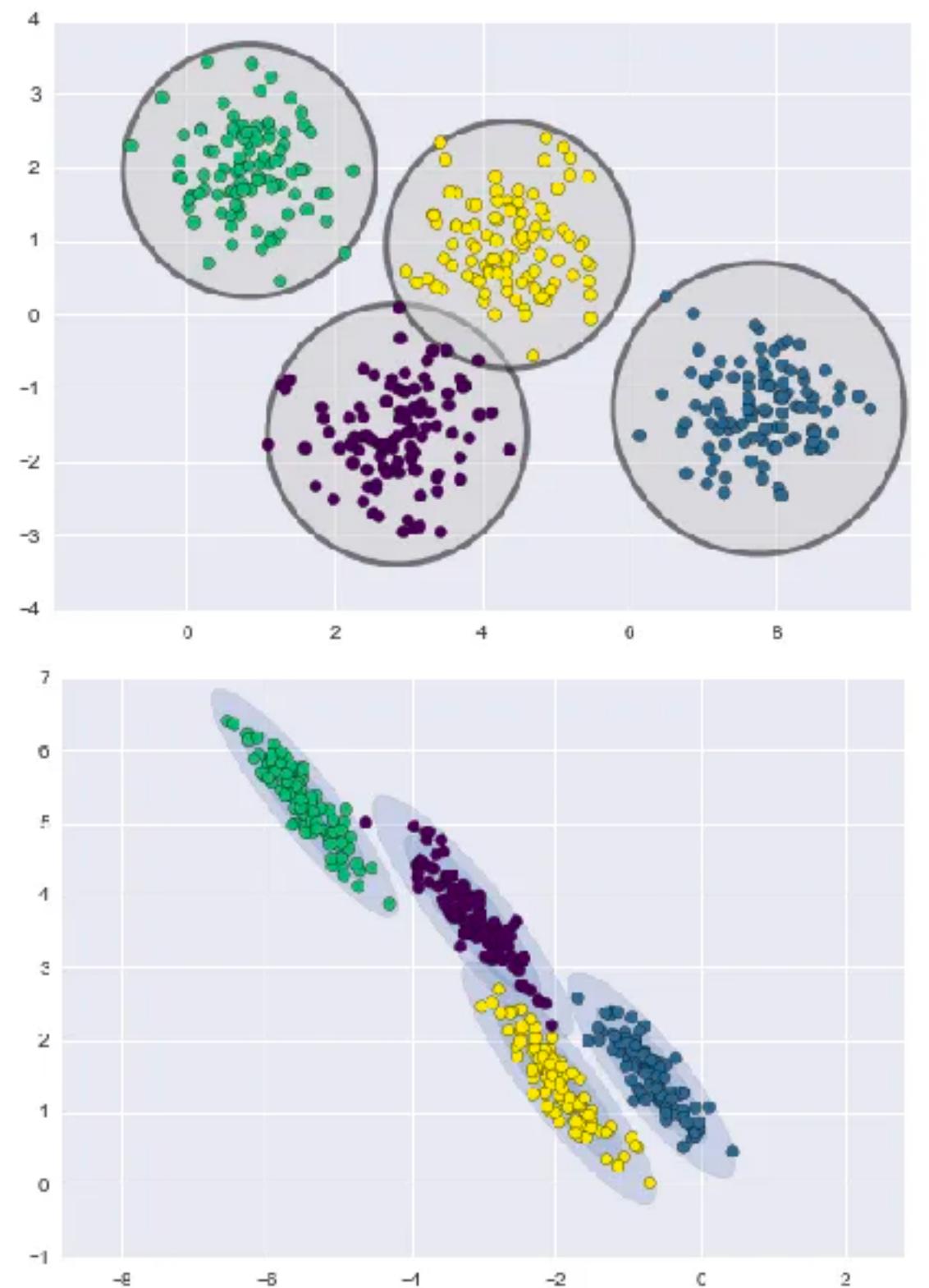


Gaussian mixture models (GMMs)

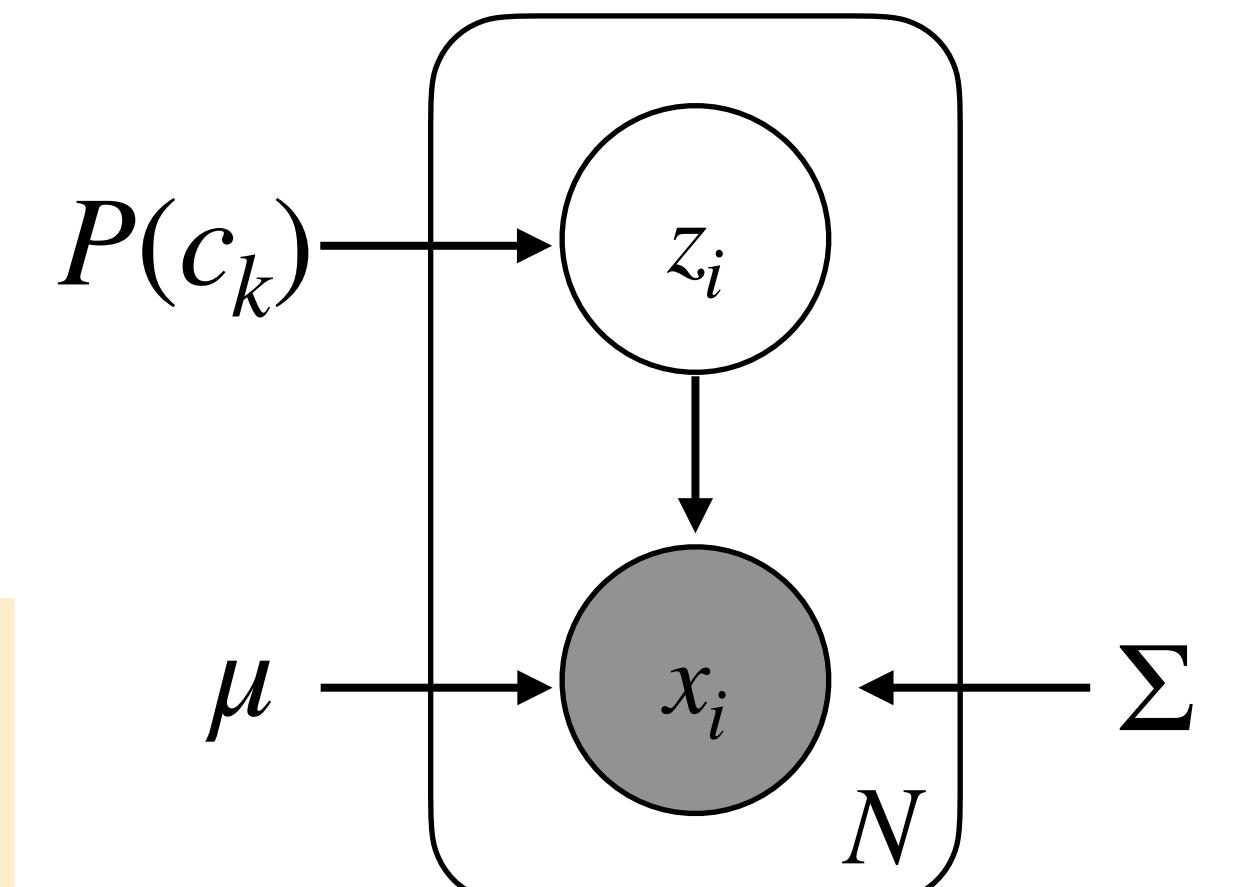
- Instead of learning a centroid (prototype), where similarity is equivalent across feature dimensions...
- ... learn a distribution for each cluster, where each feature dimension can have a different variance
 - ovals instead of spheres
- Assume data \mathbf{x}_i is generated by a latent variable z_i in the form of a Gaussian distribution with unknown means μ_k and covariance Σ_k :

$$\begin{aligned}
 & \text{Gaussian likelihood Prior} \\
 p(\mathbf{x}_i) &= \sum P(\mathbf{x}_i | z_i = k)P(z_i = k) \\
 &= \sum_k^k \mathcal{N}(\mathbf{x}_i | \mu_k, \Sigma_k)P(c_k)
 \end{aligned}$$

where $\mathcal{N}(\mathbf{x} | \mu_k, \Sigma_k) = \frac{1}{\sqrt{(2\pi)^d |\sigma|}} \exp \left(-\frac{1}{2} (\mathbf{x}_j - \mu_k)^\top \Sigma_k^{-1} (\mathbf{x}_j - \mu_k) \right)$



Graphical representation



Expectation-Maximization (EM)

- Iterative method to compute a maximum likelihood when the data depends on latent variables
- **Expectation:** Compute “expected” classes for all data points, given current parameter values

$$P(\mathbf{x}_i = c_k) = \frac{\mathcal{N}(\mathbf{x}_i | \mu_k, \Sigma_k) P(c_k)}{\sum_j^K \mathcal{N}(\mathbf{x}_i | \mu_j, \Sigma_j) P(c_j)}$$

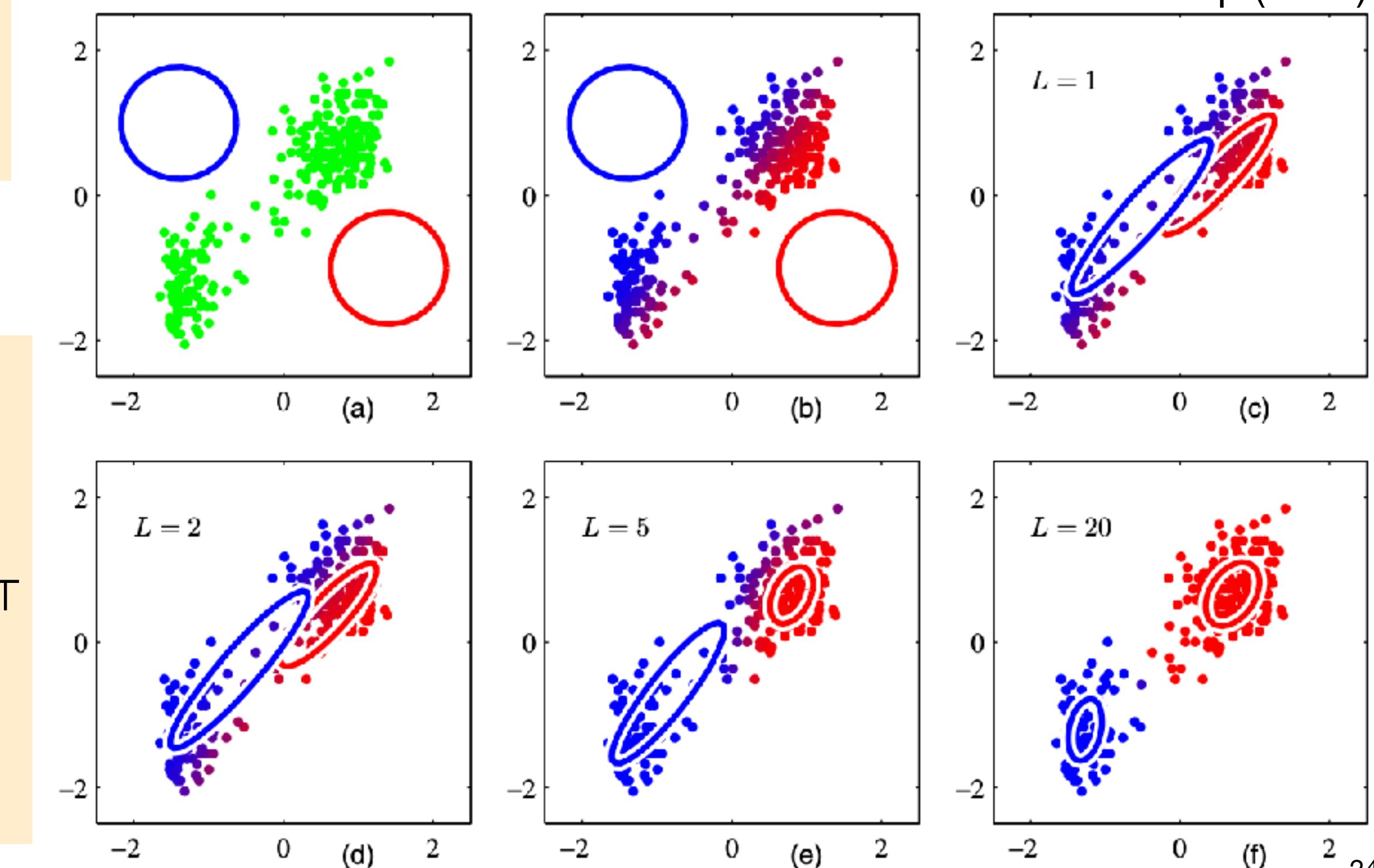
“Responsibility” of c_k for generating \mathbf{x}_i

- **Maximization:** Re-estimate parameters given current class assignments

$$\mu_k^{\text{new}} = \frac{1}{N_k} \sum_i^N P(\mathbf{x}_i = c_k) \mathbf{x}_i \quad \text{Centroid}$$

$$\Sigma_k^{\text{new}} = \frac{1}{N_k} \sum_i^N P(\mathbf{x}_i = c_k) (\mathbf{x}_i - \mu_k^{\text{new}})(\mathbf{x}_i - \mu_k^{\text{new}})^T \quad \text{Covariance}$$

$$P(c_k)^{\text{new}} = N_k / N \quad \text{Prior on classes}$$



Expectation-Maximization (EM)

What other algorithms we've covered also use similar iterative methods?

- Iterative method to compute a maximum likelihood when the data depends on latent variables
- **Expectation:** Compute “expected” classes for all data points, given current parameter values

$$P(\mathbf{x}_i = c_k) = \frac{\mathcal{N}(\mathbf{x}_i | \mu_k, \Sigma_k) P(c_k)}{\sum_j^K \mathcal{N}(\mathbf{x}_i | \mu_j, \Sigma_j) P(c_j)}$$

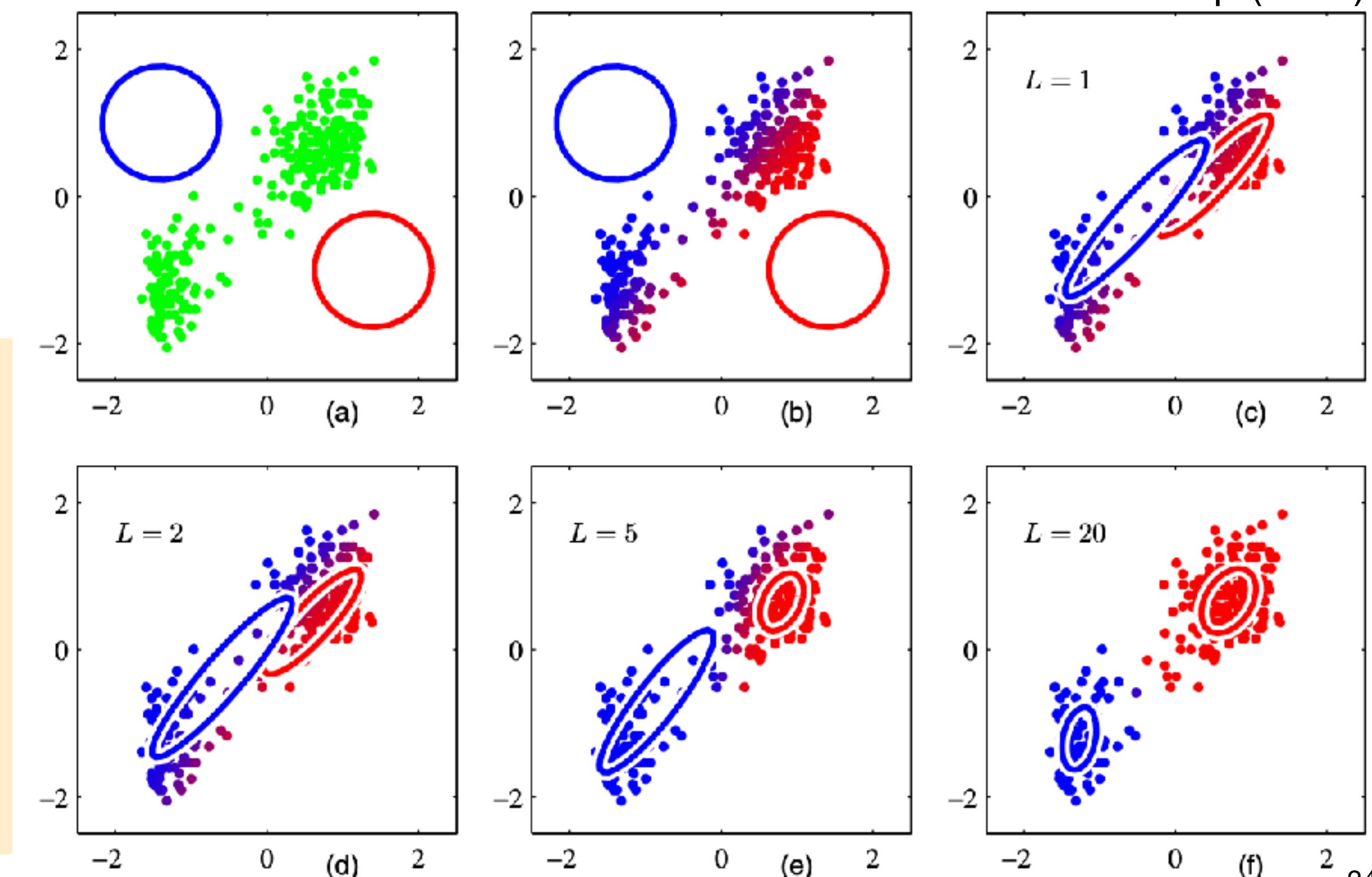
“Responsibility” of c_k for generating \mathbf{x}_i

- **Maximization:** Re-estimate parameters given current class assignments

$$\mu_k^{\text{new}} = \frac{1}{N_k} \sum_i^N P(\mathbf{x}_i = c_k) \mathbf{x}_i \quad \text{Centroid}$$

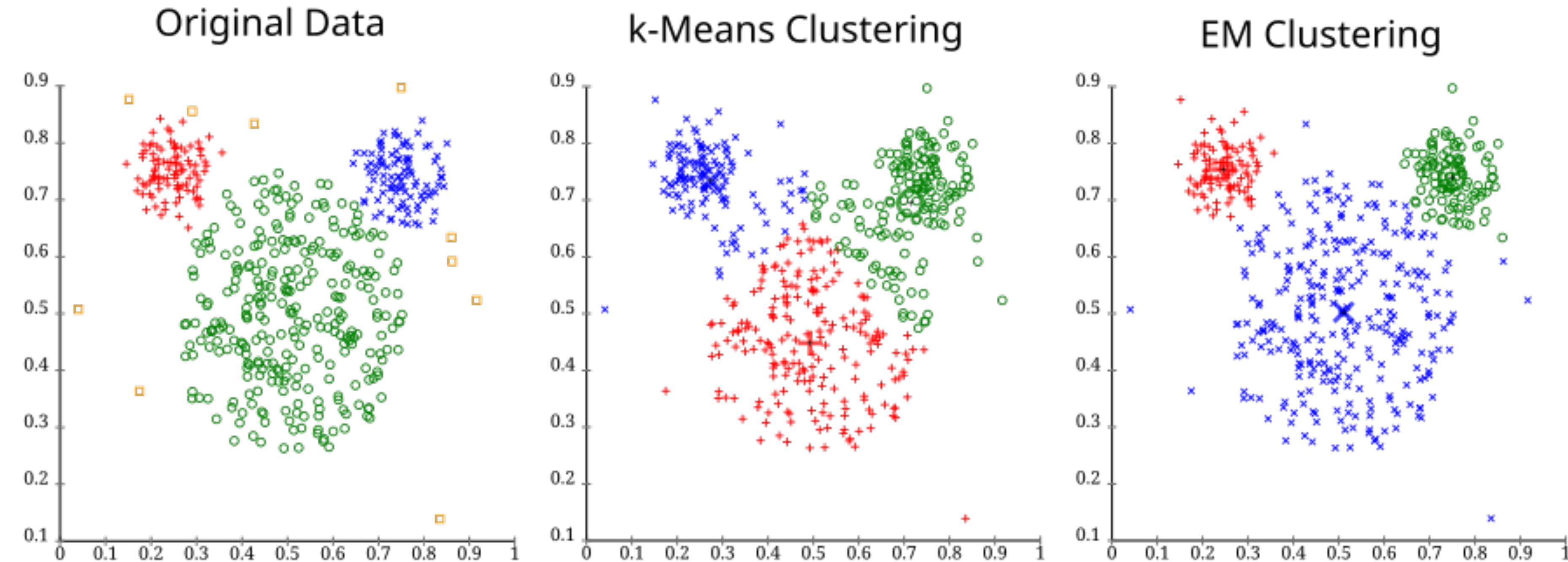
$$\Sigma_k^{\text{new}} = \frac{1}{N_k} \sum_i^N P(\mathbf{x}_i = c_k) (\mathbf{x}_i - \mu_k^{\text{new}})(\mathbf{x}_i - \mu_k^{\text{new}})^T \quad \text{Covariance}$$

$$P(c_k)^{\text{new}} = N_k / N \quad \text{Prior on classes}$$



GMMs

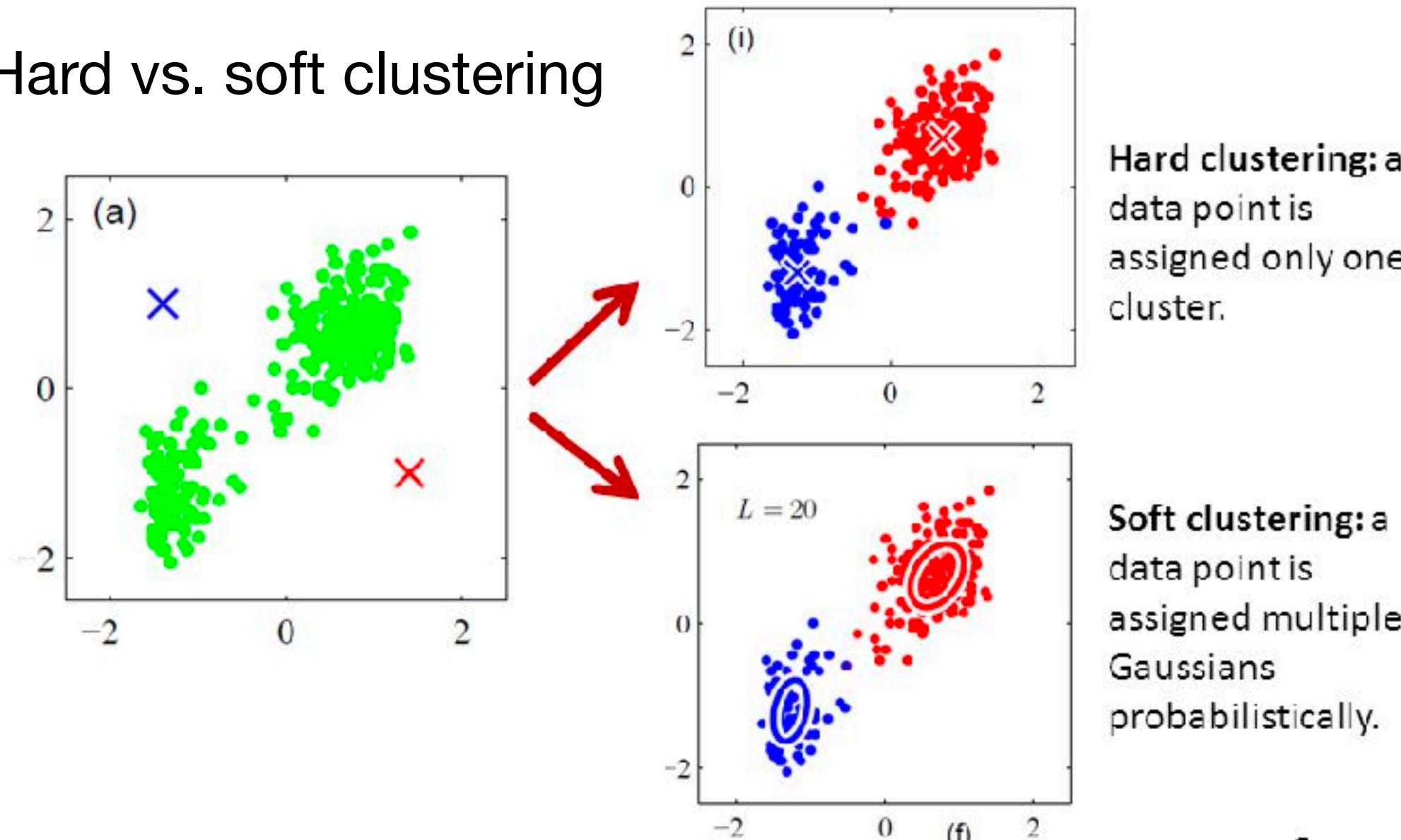
- Advantages over k -means due to
 - Learns the covariance of the data, rather than assuming it is spherical
 - Learns prior class probabilities, to account for unequal cluster sizes
- These advances could also turn into limitations, if the covariance/priors cannot be reliably estimated



Summary of unsupervised learning

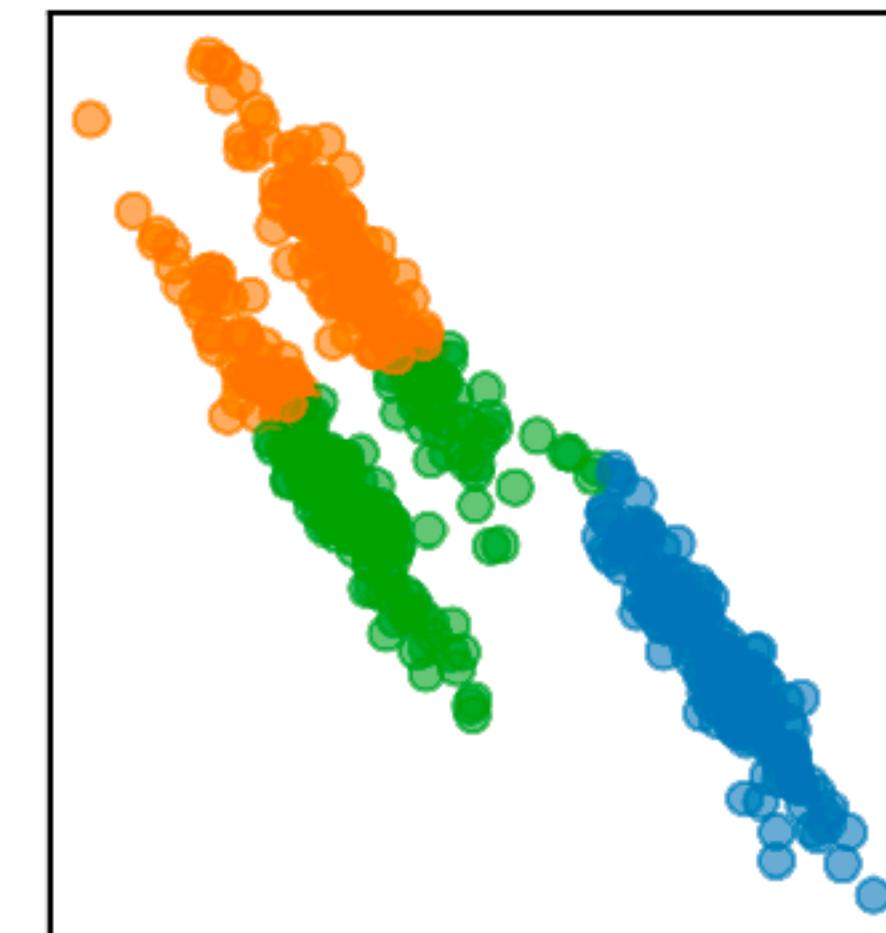
- Unsupervised learning is a clustering problem
- **k-means** performs “hard” assignment of data to clusters, with equivariant similarity across dimensions, and clusters defined by a centroid (prototype)
- **Gaussian mixture models** perform “soft” assignment, where learned covariance allows for skewed clusters, which are defined as a mixture of Gaussian densities (neither prototype or exemplar)

Hard vs. soft clustering

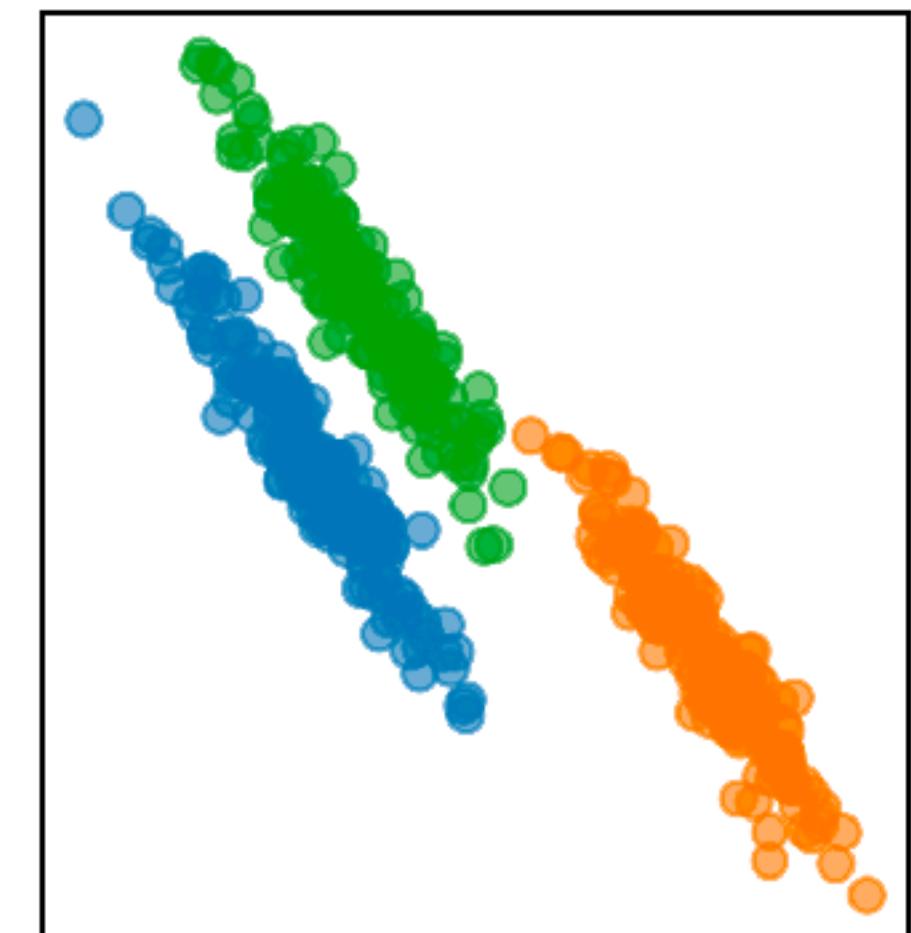


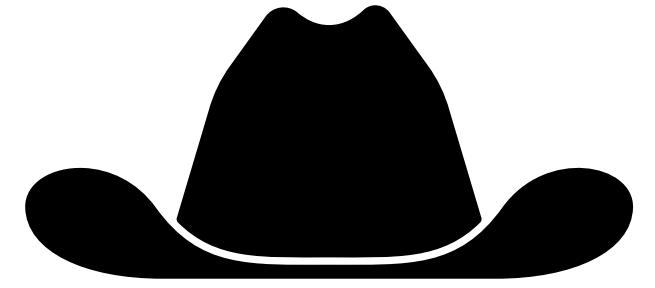
Skewed data

KMeans



GaussianMixture





Data wrangling

Beyond only implementing models, a big part of making ML work is data wrangling

- **Feature scaling**

- Min-Max normalization so feature values are in the range [0,1]

$$X' = \frac{X - X_{\min}}{X_{\max} - X_{\min}}$$

- Standardization so feature values have mean = 0 and stdev = 1

$$X' = \frac{X - \mu}{\sigma}$$

where μ is the mean and σ is the standard deviation

- Normalization vs. standardization?

- Normalization is useful when the distribution of the data is unknown or not Gaussian, since it retains the shape of the original distribution. However, it is sensitive to outliers
- Standardization is useful when the data is Gaussian (but with enough data, everything becomes Gaussian) and is less sensitive to outliers. But may change the shape of the original distribution

- **Feature engineering** by crafting new features

- e.g., # of siblings/spouses in the titanic dataset combines two separate features
- Requires some domain understanding

Assessing performance

- We need to balance both **precision** and **recall**

True Positives (TPs): 1	False Positives (FPs): 1
False Negatives (FNs): 8	True Negatives (TNs): 90

- Precision** is the proportion of items predicted TRUE that were actually TRUE

$$= \frac{TP}{TP + FP} = 1 / 1+1 = 50\%$$

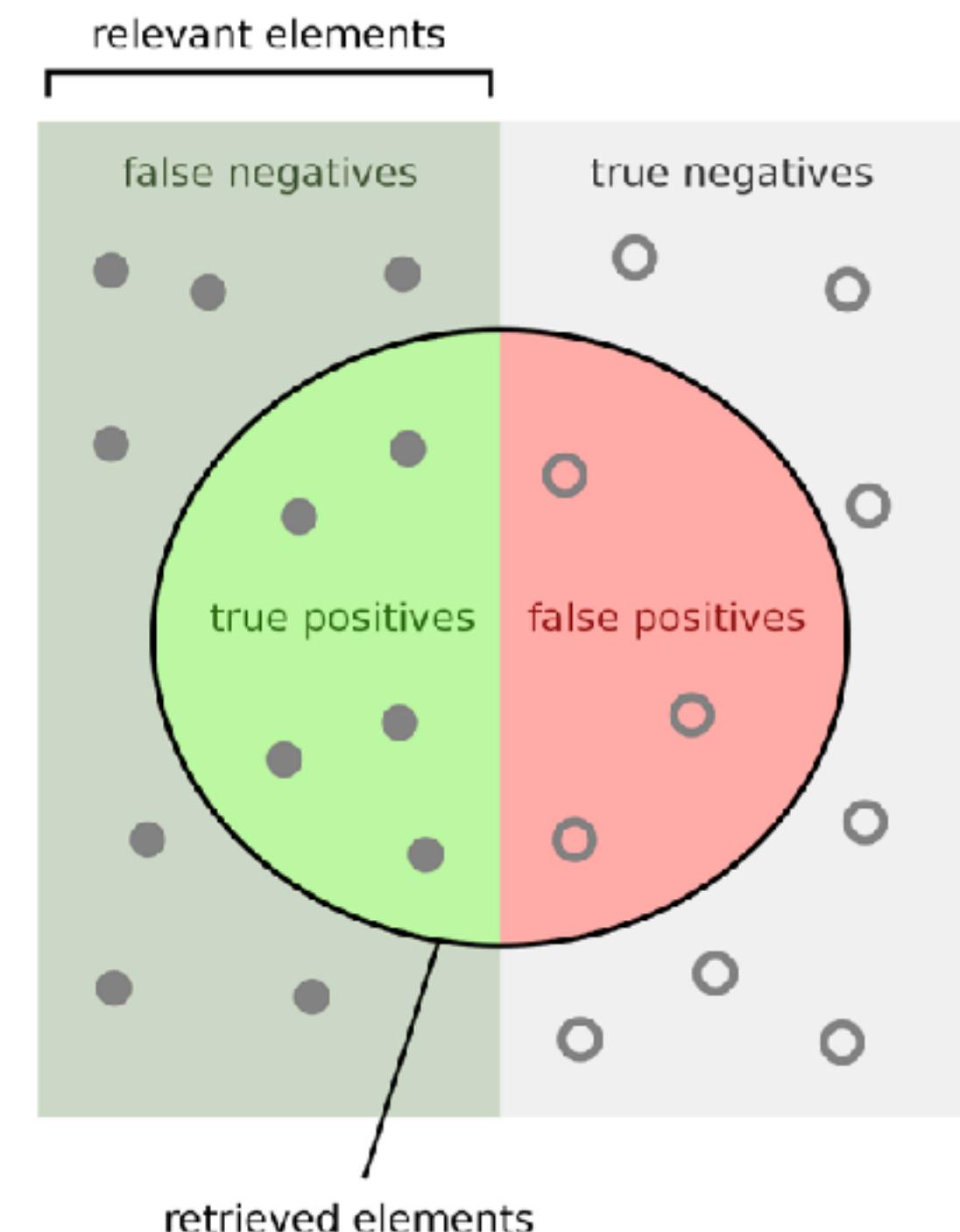
- Recall** (also known as sensitivity) is the proportion of positives that were identified correctly (i.e., labeled as TRUE)

$$= \frac{TP}{TP + FN} = 1 / 1 + 8 = 11\%$$

- Precision and recall can be a tug-of-war based on how liberal or conservative your classification algorithm is

- F1 score** is the harmonic mean of precision and recall

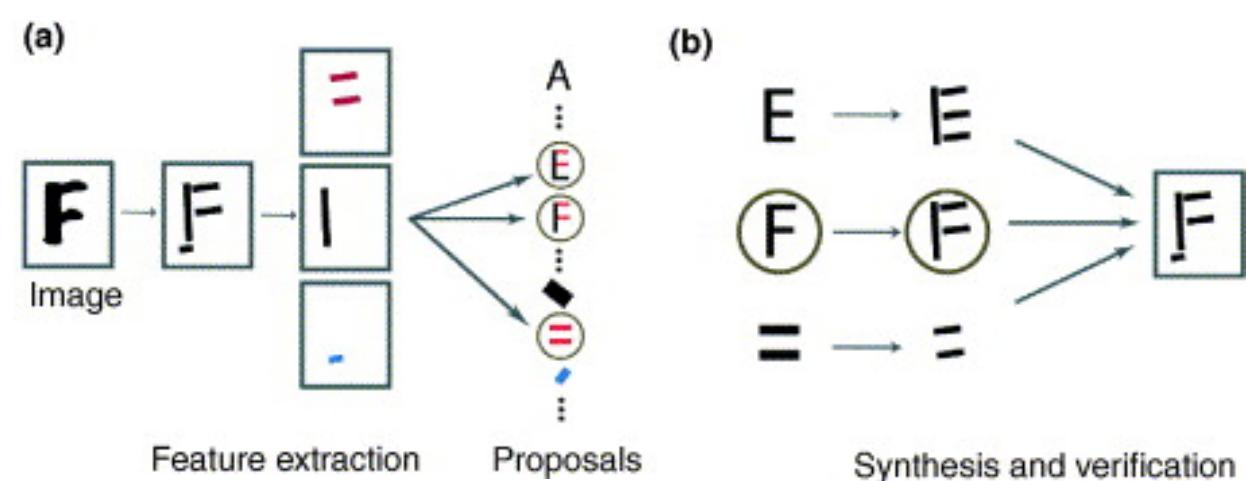
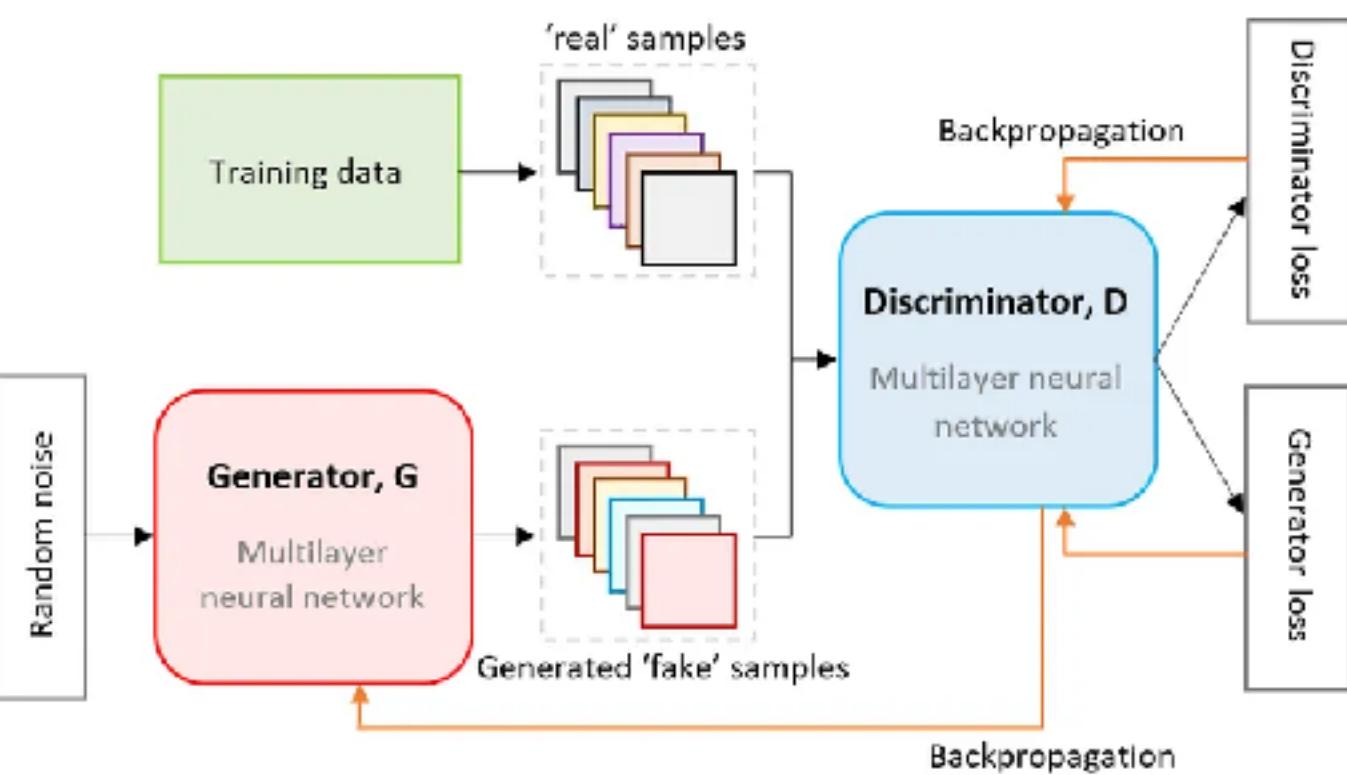
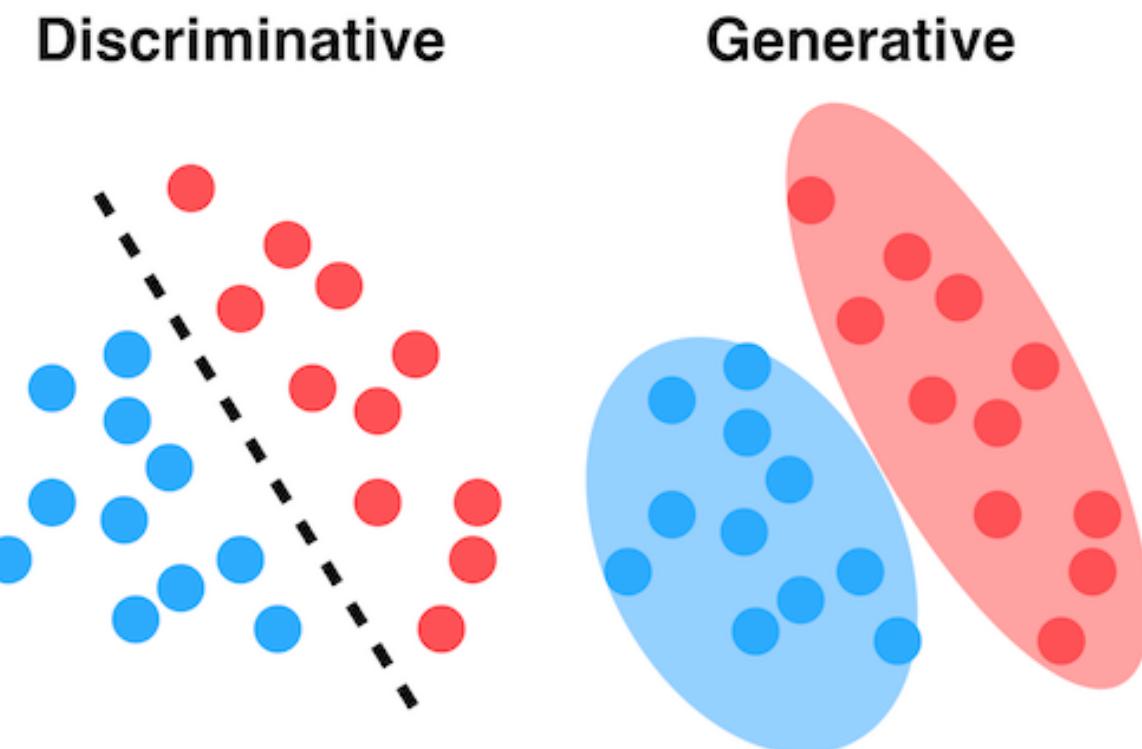
$$F1 = \frac{2}{\frac{1}{\text{Precision}} + \frac{1}{\text{Recall}}} = \frac{2 \times \text{Precision} \times \text{Recall}}{\text{Precision} + \text{Recall}} = (2 \times .5 \times .11) / (.5+.11) = 18\%$$



$$\begin{aligned} \text{Precision} &= \frac{\text{How many retrieved items are relevant?}}{\text{How many retrieved items are relevant?} + \text{How many relevant items are retrieved?}} \\ &= \frac{\text{TP}}{\text{TP} + \text{FP}} \\ \text{Recall} &= \frac{\text{How many relevant items are retrieved?}}{\text{How many relevant items are retrieved?}} \\ &= \frac{\text{TP}}{\text{TP} + \text{FN}} \\ F1 &= \frac{2 \times \text{Precision} \times \text{Recall}}{\text{Precision} + \text{Recall}} \end{aligned}$$

Discussion

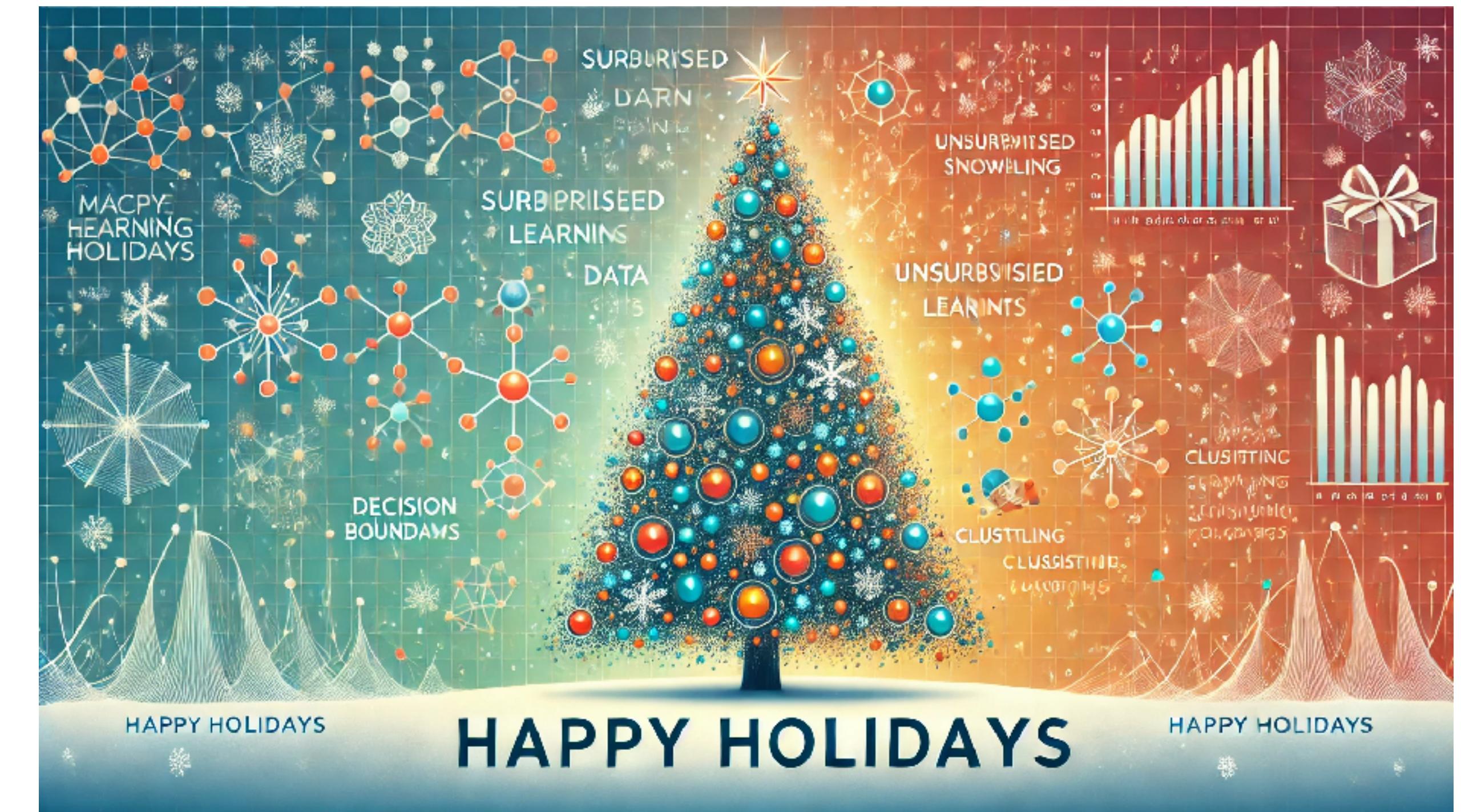
- Both supervised and unsupervised learning methods provide tools for classifying data:
 - Explicit category boundaries (decision trees, SVMs)
 - Implicit boundaries based on similarity of examples (MLPs)
 - Summary statistics of the data, based on a centroid (k-means) or a generative distribution (Naïve Bayes, GMM)
- **Discriminative** models simply learn to recognize the category labels (decision trees, SVMs, k-means)
- **Generative** models (Naïve Bayes, GMM) learn the data distribution and can be used to generate new datapoints consistent with each category
- Many ML methods combine both (e.g., GANs, Bayesian adversarial networks)
 - Discriminative models are cheap to learn, but require a lot of data
 - Generative models are more computationally costly, but can generate additional training data
 - Discriminative model provides an additional training signal to generative model, while generative model can simulate more training data (similar to model-based planning in RL)
- “Analysis by synthesis” (Yuille & Kersten, 2006) suggests humans do something similar, through an interaction between top-down generative processes and bottom-up discrimination



Bring your laptops for the tutorial on Friday

- We will provide 1 supervised and 1 unsupervised classification dataset
- Given the training data, implement one model of your choice
- We will provide code examples in Python and R for each model covered today
- Then, test your models on the test set. Best test performance on each dataset wins a prize!
- We will use F1 score as the performance metric

Next week



See you on Jan 14th for Function learning

Function Learning

