

# Machine Learning and Pattern Recognition A High Level Overview

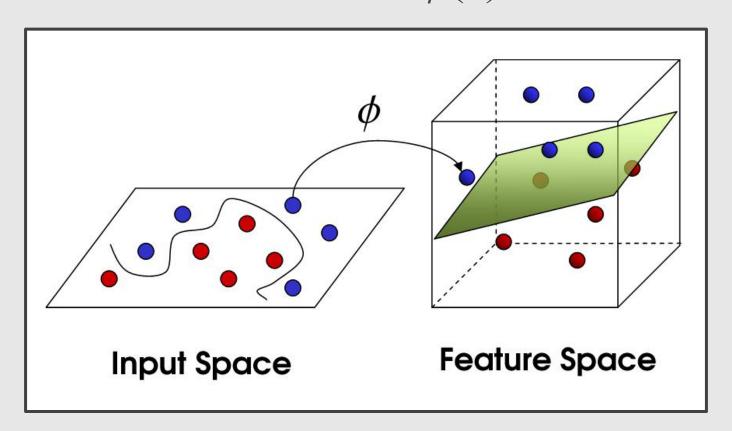
#### Prof. Anderson Rocha

(Main bulk of slides kindly provided by **Prof. Sandra Avila**)
Institute of Computing (IC/Unicamp)

# Non-linear SVMs: Feature spaces

General idea: the original feature space can always be mapped to some higher-dimensional feature space where the training set is separable.

$$\Phi: x \to \varphi(x)$$



# Kernel Trick

- The linear classifier relies on inner product between vector  $K(x_i, x_j) = x_i^T x_j$
- If every datapoint is mapped into high-dimensional space via some transformation  $\Phi: x \to \varphi(x)$ , the inner product becomes:

$$K(x_i, x_j) = \varphi(x_i)^{\mathrm{T}} \varphi(x_j)$$

• A kernel function is a function that is equivalent to an inner product in some feature space.

Suppose you want to apply a 2<sup>nd</sup> degree polynomial transformation to a 2-dimensional training set, then train a linear SVM classifier on the transformed training set.

$$\varphi(x) = \varphi((x_1 \ x_2)) = (x_1^2 \ \sqrt{2x_1x_2} \ x_2^2)$$

The transformed vector is 3-dimensional instead of 2-dimensional.

Let's look at what happens to a couple 2-dimensional vectors **a** and **b** if we apply this 2<sup>nd</sup> degree polynomial mapping then compute the dot product of the transformed vectors.

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$$\varphi(\mathbf{a})^{\mathrm{T}}\varphi(\mathbf{b}) = (a_1^2 \sqrt{2}a_1a_2 a_2^2)^{\mathrm{T}}(b_1^2 \sqrt{2}b_1b_2 b_2^2) =$$

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$$= a_{1}^{2}b_{1}^{2} + 2a_{1}a_{2}b_{1}b_{2} + a_{2}^{2}b_{2}^{2} =$$

$$= (a_{1}b_{1} + a_{2}b_{2})^{2} =$$

$$= ((a_{1} a_{2})^{\mathrm{T}}(b_{1} b_{2}))^{2} =$$

$$= (\mathbf{a}^{\mathrm{T}} \cdot \mathbf{b})^{2}$$

The dot product of the transformed vectors is equal to the square of the dot product of the original vectors:  $\varphi(\mathbf{a})^T \varphi(\mathbf{b}) = (\mathbf{a}^T \cdot \mathbf{b})^2$ 

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So you don't actually need to transform the training instances at all: just replace the dot product by its square.

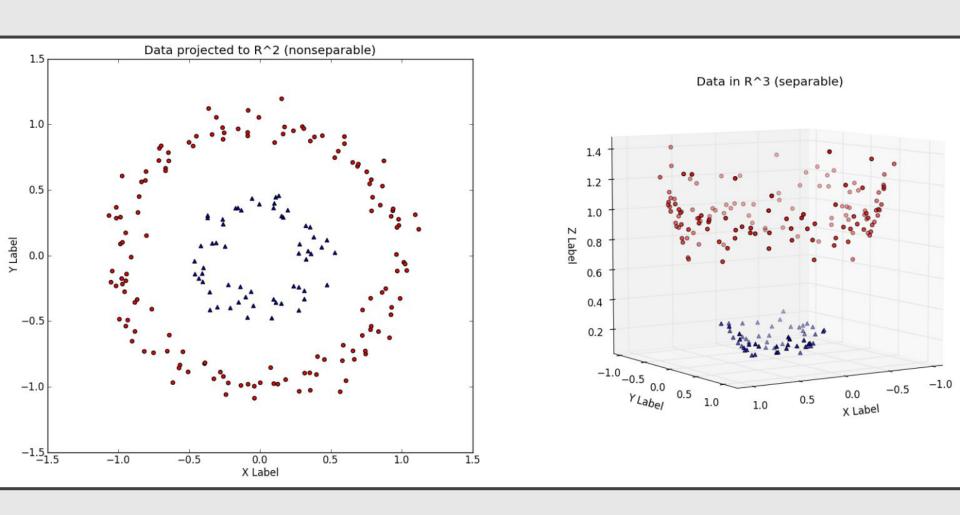
This is the essence of the kernel trick.

In Machine Learning, a *kernel* is a function capable of computing the dot product  $\varphi(\mathbf{a})^T \varphi(\mathbf{b})$  based only on the original vectors  $\mathbf{a}$  and  $\mathbf{b}$ , without having to compute (or even to know about) the transformation  $\varphi$ .

# Kernel Functions

- Linear SVM = Kernel SVM:  $K(x_i, x_j) = x_i^T x_j$
- Nonlinear SVM:  $K(x_i, x_j) = \varphi(x_i) \cdot \varphi(x_j)$

- Gaussian kernel:  $K(x_i, x_j) = \exp(-\|x_i x_j\|^2/(2\sigma^2))$
- Polynomial kernel:  $K(x_i, x_j) = (x_i \cdot x_j + 1)^d$ , d degree
- Chi-square kernel, histogram intersection kernel, string kernel, ....



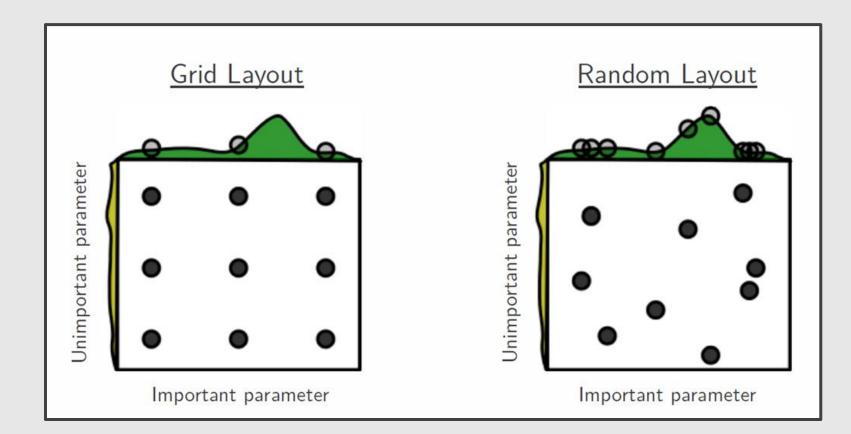
# Important Parameters

Important parameters having higher impact on model performance, "kernel", "gamma" and "C".

The parameters can be tuned using grid-search.



# Grid Search



#### References

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#### **Machine Learning Books**

- Hands-On Machine Learning with Scikit-Learn and TensorFlow, Chap. 5
- Pattern Recognition and Machine Learning, Chap. 6 & 7

#### **Machine Learning Courses**

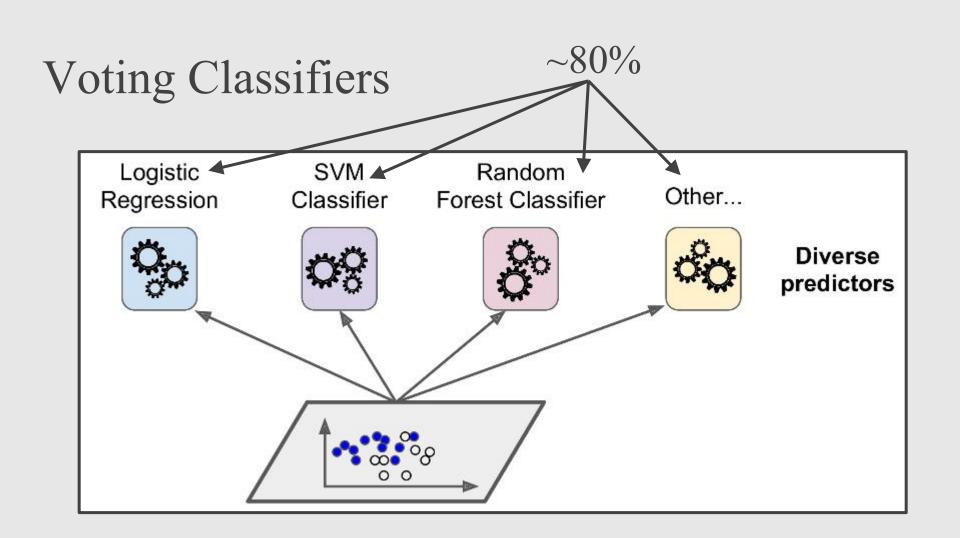
- <a href="https://www.coursera.org/learn/machine-learning">https://www.coursera.org/learn/machine-learning</a>, Week 7
- <a href="http://cs229.stanford.edu/syllabus.html">http://cs229.stanford.edu/notes/cs229-notes3.pdf</a>



# Ensemble Learning and Random Forests Machine Learning and Pattern Recognition

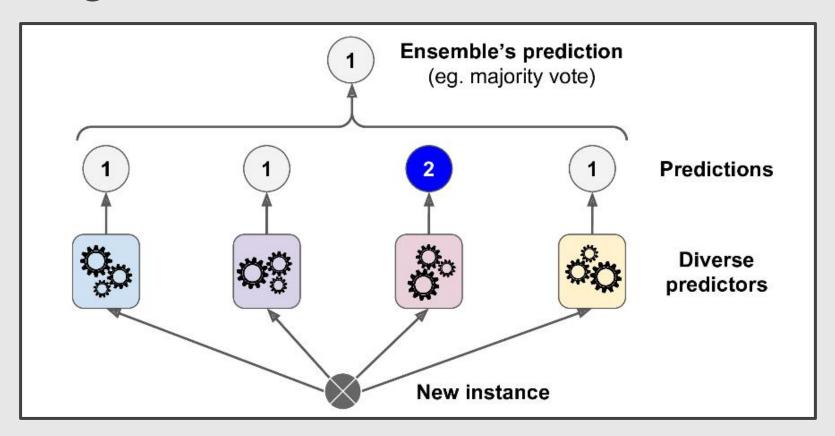
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# Voting Classifiers

#### Hard voting classifier

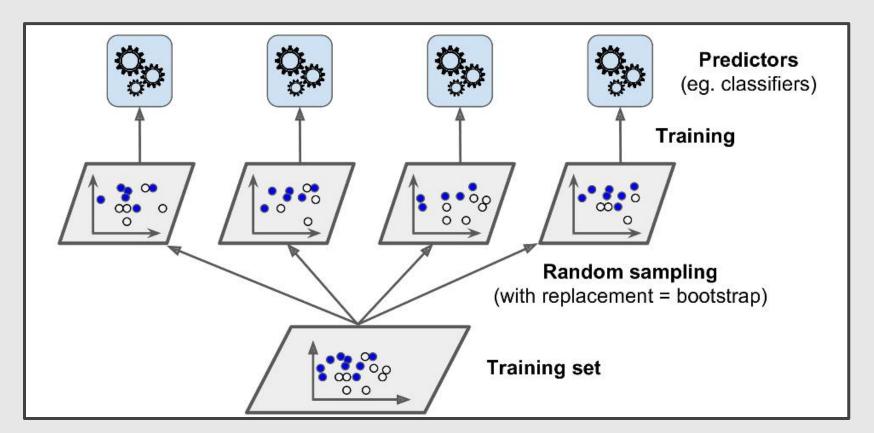


# Voting Classifiers

- Voting classifier often achieves a higher accuracy than the best classifier in the ensemble.
- Even if each classifier is a weak learner (meaning it does only slightly better than random guessing), the ensemble can still be a strong learner (achieving high accuracy).

• Use the same training algorithm for every predictor, but to train them on different random subsets of the training set.

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- Bagging (short for Bootstrap Aggregating): sampling is performed with replacement.
- Pasting: sampling is performed without replacement



- Once all predictors are trained, the ensemble can make a prediction for a new instance by simply aggregating the predictions of all predictors.
- Bagging and Pasting scale very well.

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- Bagging and Pasting scale very well.
- Out-of-bag evaluation: m% for training and (1-m)% for test

# Decision Tree & Random Forest

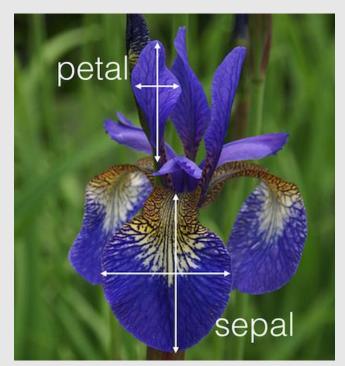
# Decision Tree & Random Forest

 Decision Trees are versatile Machine Learning algorithms that can perform both classification and regression tasks, and even multi-output tasks.

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- Random Forest is an ensemble of Decision Trees, generally trained using the Bagging method (or sometimes Pasting).

#### Decision Tree: Iris Dataset



http://sebastianraschka.com/Articles/2014\_python\_lda.html

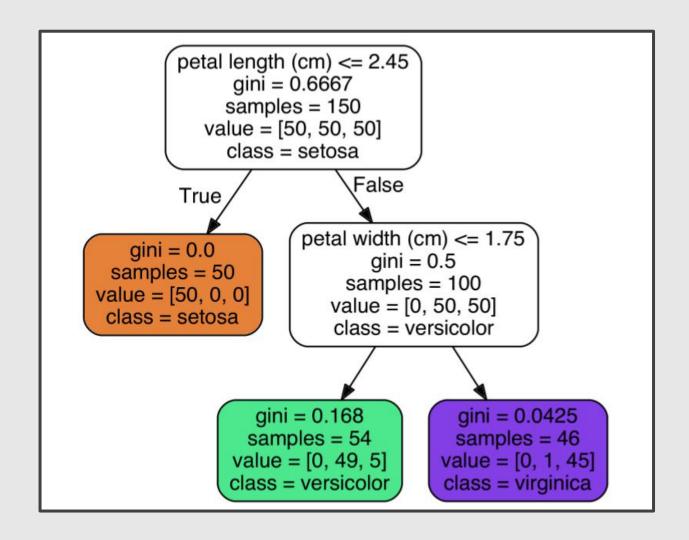
150 iris flowers from three different species.

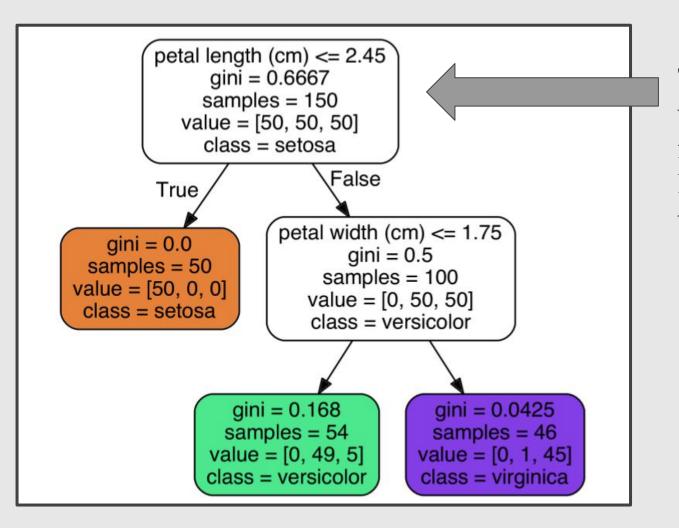
The three classes in the Iris dataset:

- 1. Iris-setosa (n=50)
- 2. Iris-versicolor (n=50)
- 3. Iris-virginica (*n*=50)

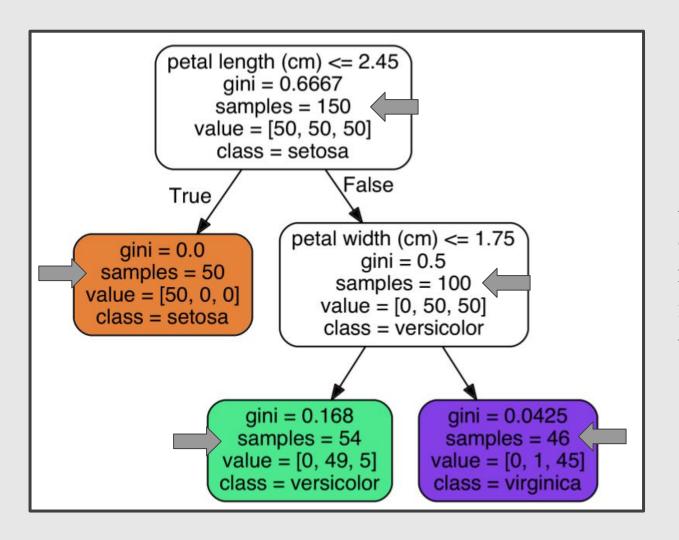
The four features of the Iris dataset:

- 1. sepal length in cm
- 2. sepal width in cm
- 3. petal length in cm
- 4. petal width in cm

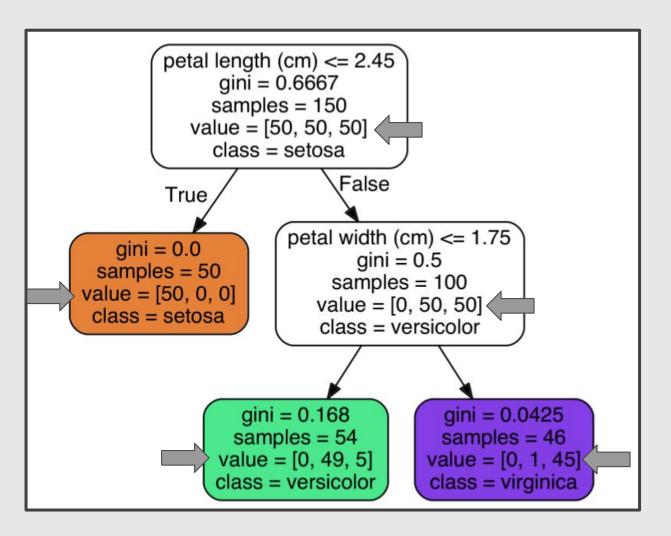




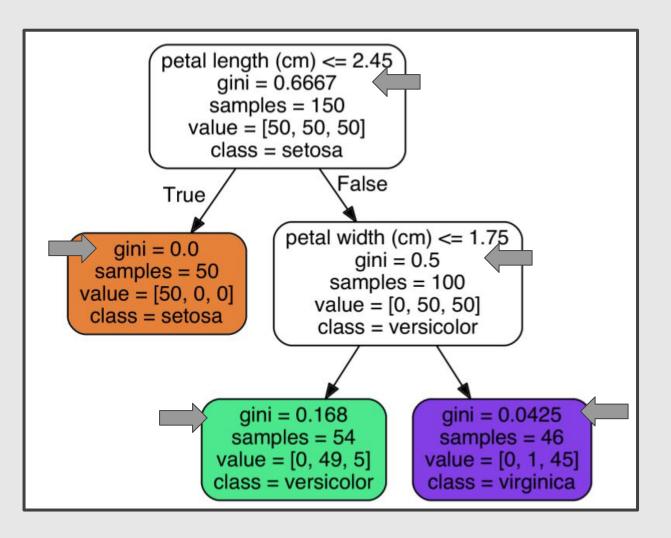
This node asks whether the flower's petal length is smaller than 2.45 cm



A node's samples attribute counts how many training instances it applies to.



A node's value attribute tells you how many training instances of each class this node applies to.



A **node's gini** attribute measures its impurity.

"pure" (gini=0): all training instances belong to the same class.

petal length (cm) <= 2.45 gini = 0.6667samples = 150

For example, the depth 2 left node has a gini score equal to  $1 - (0/54)^2 - (49/54)^2 - (5/54)^2 \approx 0.168$ .

$$G_i = 1 - \sum p_{i,k}^2$$

 $p_{i,k}$  is the ratio of class k instances among the training instances in the ith node

gini = 0.168samples = 54value = [0, 49, 5]class = versicolor

samples = 46value = [0, 1, 45]class = virginica

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•0): all nces same

• Classification And Regression Tree (CART) algorithm.

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- The idea is really quite simple: the algorithm first splits the training set in two subsets using a single feature k and a threshold  $t_k$  (e.g. "petal length  $\leq 2.45$  cm").

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- The idea is really quite simple: the algorithm first splits the training set in two subsets using a single feature k and a threshold  $t_k$  (e.g. "petal length  $\leq 2.45$  cm").
- How does it choose k and  $t_k$ ? The answer is that it searches for the pair  $(k, t_k)$  that produces the purest subsets (weighted by their size).

$$J(k, t_k) = \frac{m_{\rm left}}{m} G_{\rm left} + \frac{m_{\rm right}}{m} G_{\rm right}$$
 where 
$$\begin{cases} G_{\rm left/right} \text{ measures the impurity of the left/right subset,} \\ m_{\rm left/right} \text{ is the number of instances in the left/right subset.} \end{cases}$$

CART cost function for classification

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#### CART cost function for classification

It stops recursing once it reaches the maximum depth (hyperparameter), or if it cannot find a split that will reduce impurity.

To be continued ...

#### References

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#### **Machine Learning Books**

- Hands-On Machine Learning with Scikit-Learn and TensorFlow, Chap. 6 & 7
- Pattern Recognition and Machine Learning, Chap. 14
- Pattern Classification, Chap 8 & 9 (Sec. 9.5)