Classification

(Slides largely based on Prof. Sandra Avila's Machine Learning Course)

Prof. Rosa Paccotacya Yanque

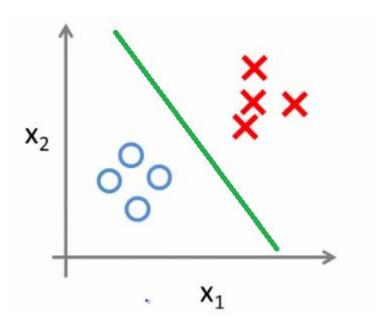
Agenda

- Logistic Regression
- Ensemble Learning
- Support Vector Machines (SVMs)
- Decision Tree
- Random Forest

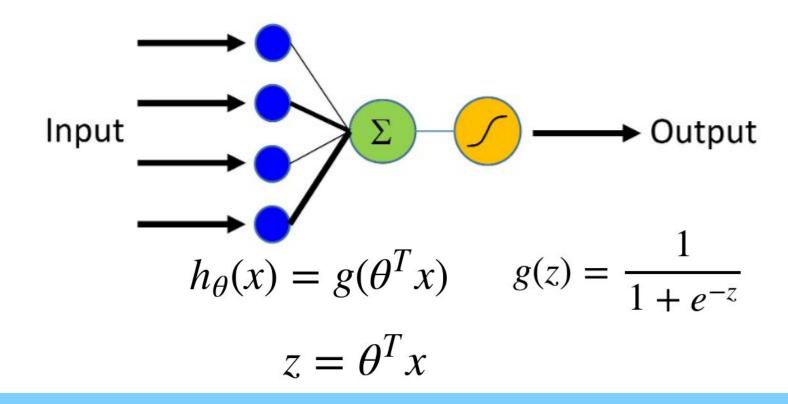
Logistic Regression

Binary Classification

- Transactions(fraudulent or no)
- People with disease or no
- Tumors (benign or not)
- E-mail (spam or not)



Model



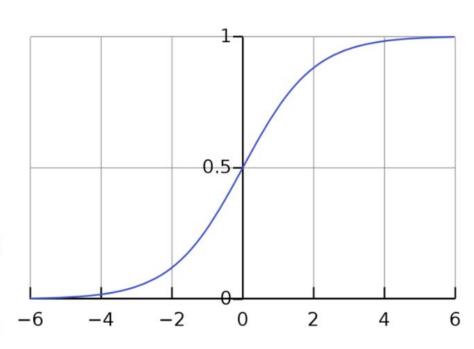
Sigmoid or Logistic Function

$$g(z) = \frac{1}{1 + e^{-z}}$$

$$z = \theta^T x$$

$$z=0, e^0=1\Rightarrow g(z)=1/2 \ z o\infty, e^{-\infty} o 0\Rightarrow g(z)=1$$

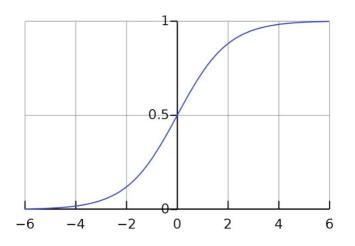
$$z
ightharpoonup -\infty, e^{\infty}
ightharpoonup \infty \Rightarrow g(z) = 0$$
 ⁻⁶



Interpretation

$$h_{ heta}(x) \geq 0.5
ightarrow y = 1 \ h_{ heta}(x) < 0.5
ightarrow y = 0$$

$$h_{ heta}(x) = g(heta^T x) \geq 0.5 \ when \ heta^T x \geq 0$$



$$egin{aligned} heta^T x &\geq 0 \Rightarrow y = 1 \ heta^T x &< 0 \Rightarrow y = 0 \end{aligned}$$

$$h_{\theta}(x) = g(\theta_0 + \theta_1 x_1 + \theta_2 x_2)$$

Predict "
$$y = 1$$
" if $-3 + x_1 + x_2 \ge 0$

Training set: $\{(x^{(1)}, y^{(1)}), (x^{(2)}, y^{(2)}), \cdots, (x^{(m)}, y^{(m)})\}$

$$x \in \begin{bmatrix} x_0 \\ x_1 \\ \dots \\ x_n \end{bmatrix}$$

m examples
$$x \in \begin{bmatrix} x_0 \\ x_1 \\ \dots \\ x_n \end{bmatrix}$$
 $x_0 = 1, y \in \{0, 1\}$

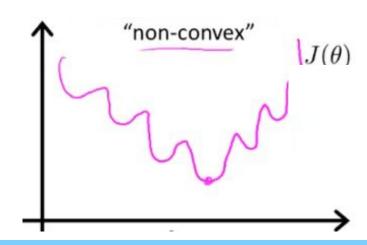
$$h_{\theta}(x) = \frac{1}{1 + e^{-\theta^T x}}$$

How to choose parameters θ ?

$$Cost(h_{\theta}(x^{(i)}), y^{(i)}) = \frac{1}{2} \left(h_{\theta}(x^{(i)}) - y^{(i)} \right)^{2}$$

Linear regression: $J(\theta) = \frac{1}{m} \sum_{i=1}^{m} \frac{1}{2} \left(h_{\theta}(x^{(i)}) - y^{(i)} \right)^2$

Cost Function

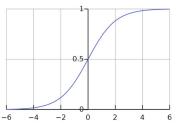


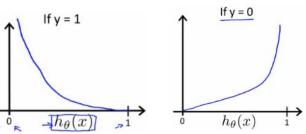
LR COST FUNCTION

$$J(heta) = rac{1}{m} \sum_{i=1}^m \mathrm{Cost}(h_{ heta}(x^{(i)}), y^{(i)})$$

$$\operatorname{Cost}(h_{\theta}(x), y) = -\log(h_{\theta}(x))$$
 if $y = 1$
 $\operatorname{Cost}(h_{\theta}(x), y) = -\log(1 - h_{\theta}(x))$ if $y = 0$

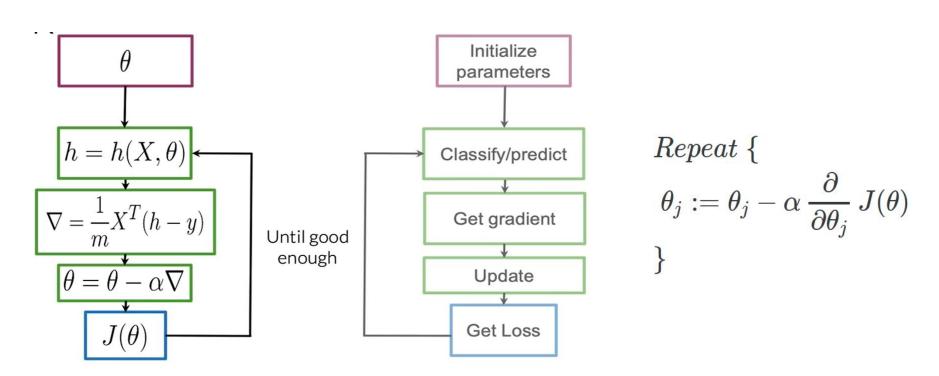
$$\operatorname{Cost}(h_{\theta}(x), y) = -\log(1 - h_{\theta}(x))$$





$$Cost(h_{\theta}(x), y) = -y \log(h_{\theta}(x)) - (1 - y) \log(1 - h_{\theta}(x))$$

Training:Gradient Descent



Multiclass Classification

- Email tagging: Work, Friends, Family
- Skin Lesion: Melanoma, Carcinoma, Nevus, Keratosis
- Video: Pornography, Violence, Gore scenes, Child abuse

Classification

Email tagging: Work, Friends, Family

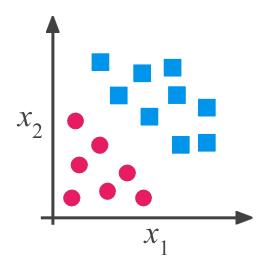
$$y = 1$$
 $y = 2$ $y = 3$

Skin Lesion: Melanoma, Carcinoma, Nevus, Keratosis

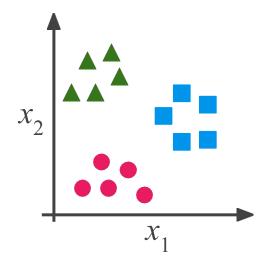
$$y = 1$$
 $y = 2$ $y = 3$ $y = 4$

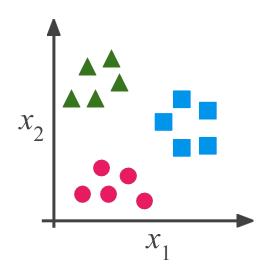
Video: Pornography, Violence, Gore scenes, Child abuse

Binary Classification



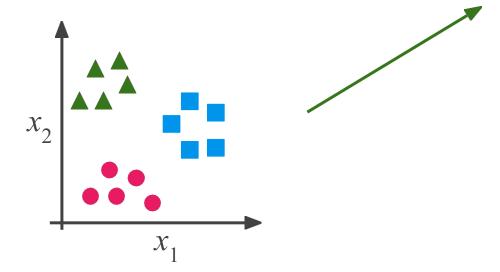
Multi-class Classification

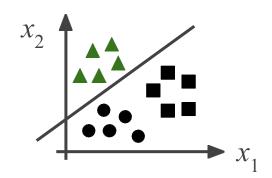




Class 1: ▲

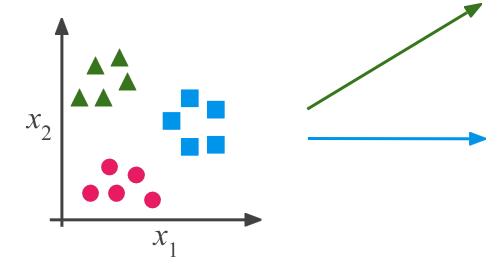
Class 2:

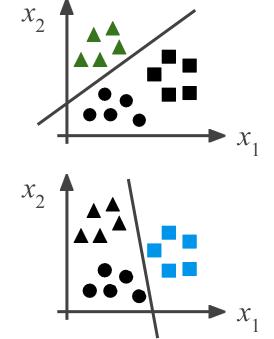




Class 1: ▲

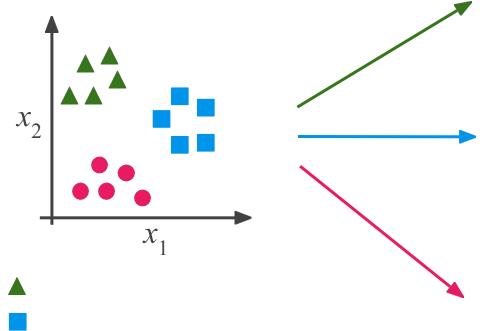
Class 2:

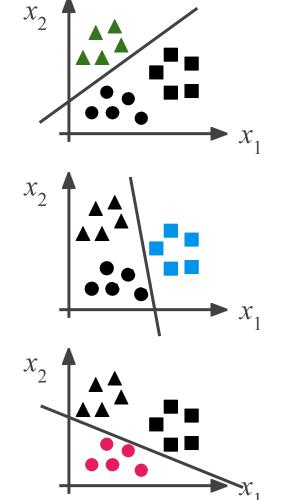




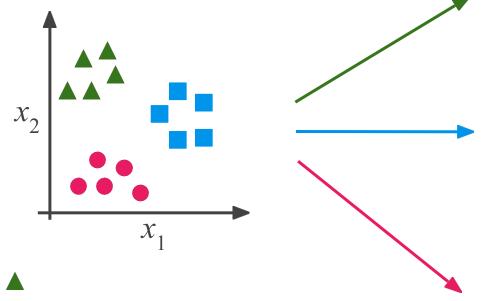
Class 1:

Class 2:





Class 1: ▲ Class 2: ■

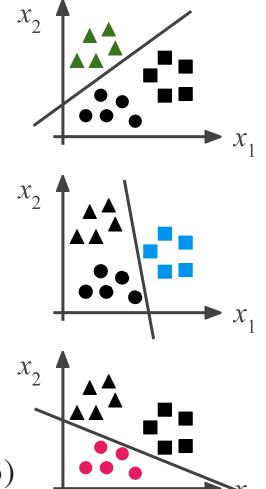


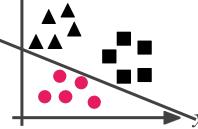
Class 1:

Class 2:

$$h_{\theta}^{(i)}(x) = P + y \quad i \mid x; \theta) \quad (i=1,2,3)$$

$$(i=1,2,3)$$





Train a logistic regression classifier $h_{\theta}^{(i)}(x)$ for each class i to predict the probability that y = i.

One a new input x, to make a prediction, pick the class i that maximizes

$$\max_{i} h_{\theta}^{(i)}(x)$$

References

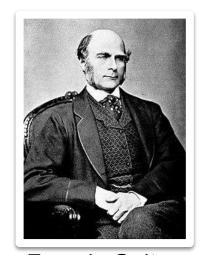
Machine Learning Books

- Hands-On Machine Learning with Scikit-Learn and TensorFlow, Chap. 4
- Pattern Recognition and Machine Learning, Chap. 4

Machine Learning Courses

- https://www.coursera.org/learn/machine-learning, Week 3
- Logistic Regression The Math of Intelligence (Week 2): https://youtu.be/D8alok2P468
- http://cs229.stanford.edu/notes/cs229-notes1.pdf

Ensemble Learning



Francis Galton (1822-1909)

Animal's weight?





~800 people 1,197 kg

1,207 kg

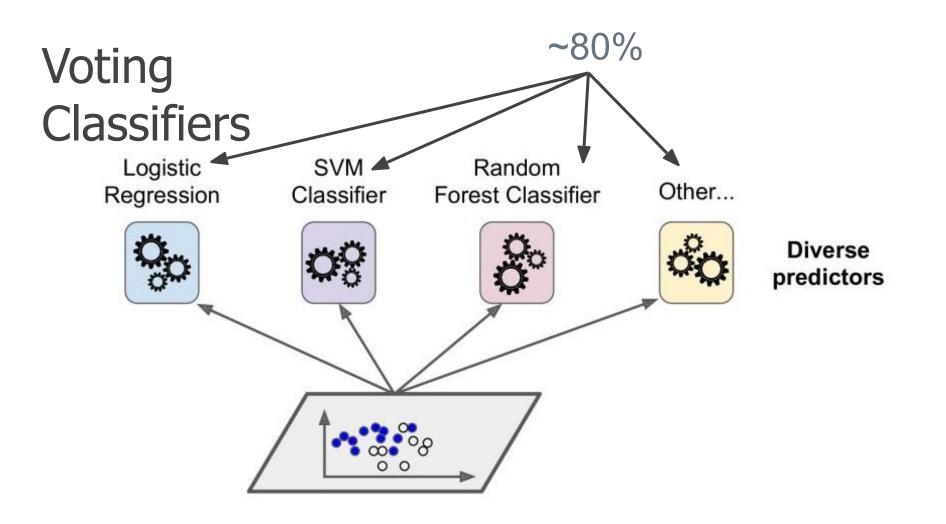


Wisdom of the Crowd

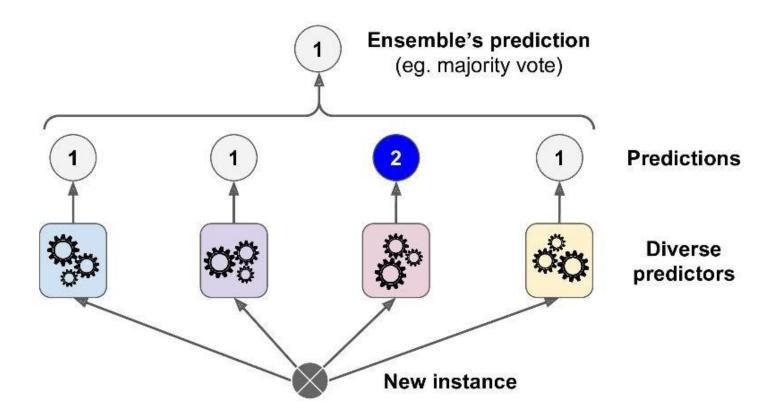


Ensemble Learning

 Multiple learning algorithms to obtain better <u>predictive</u> <u>performance</u> than could be obtained from any learning algorithms individually.



Hard/Soft voting classifier



 Voting classifier often achieves a higher accuracy than the best classifier in the ensemble.

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• Even if each classifier is a weak learner, the ensemble can still be a strong learner, provided there are a sufficient number of weak learners and they are sufficiently diverse.

 Ensemble methods work best when the predictors are as independent from one another as possible.

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 One way to get diverse classifiers is to train them using very different algorithms: this increases the chance that they will make very different types of errors, improving the ensemble's accuracy.

```
from sklearn.ensemble import RandomForestClassifier
from sklearn.ensemble import VotingClassifier
from sklearn.linear model import LogisticRegression
from sklearn.svm import SVC
log clf = LogisticRegression()
rnd clf = RandomForestClassifier()
svm clf = SVC()
voting clf = VotingClassifier(
        estimators=[('lr', log clf), ('rf', rnd clf), ('svc', svm clf)],
                     voting='hard'
voting clf.fit(X train, y train)
```

LogisticRegression 0.864

RandomForestClassifier 0.896

Voting Classifiers

```
SVC 0.888
from sklearn.ensemble import RandomForestClassif
                                                   VotingClassifier 0.904
from sklearn.ensemble import VotingClassifier
from sklearn.linear model import LogisticRegression
from sklearn.svm import SVC
log clf = LogisticRegression()
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Ensemble Methods

- Bagging (and Pasting)
- Boosting
- Stacking

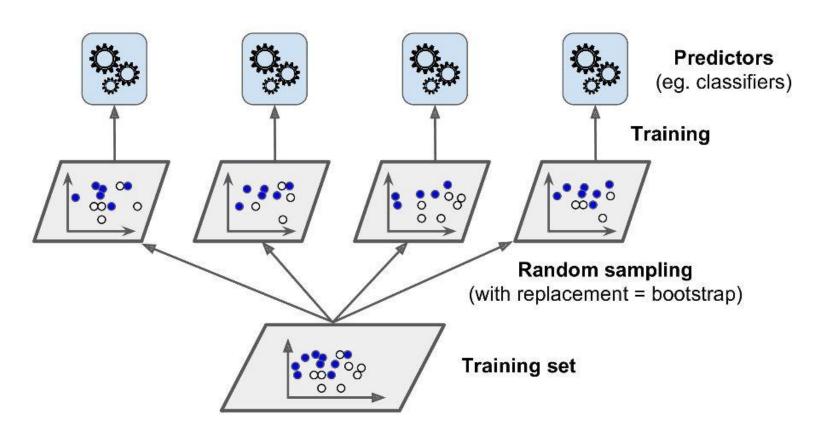
Bagging and Pasting

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

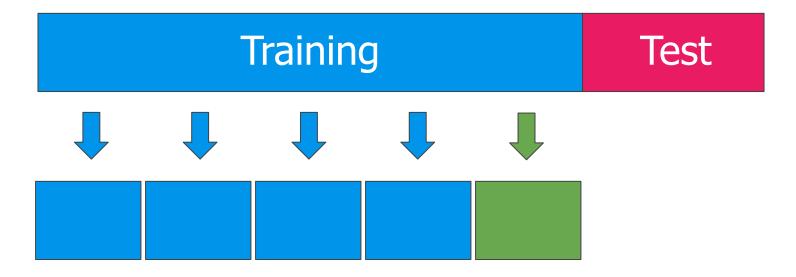
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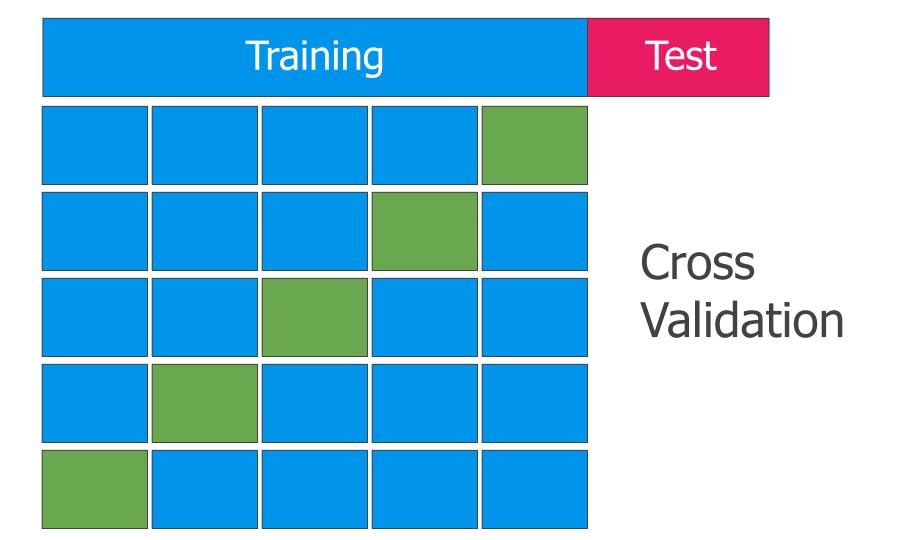
 Bagging (short for Bootstrap Aggregating): sampling is performed with replacement.

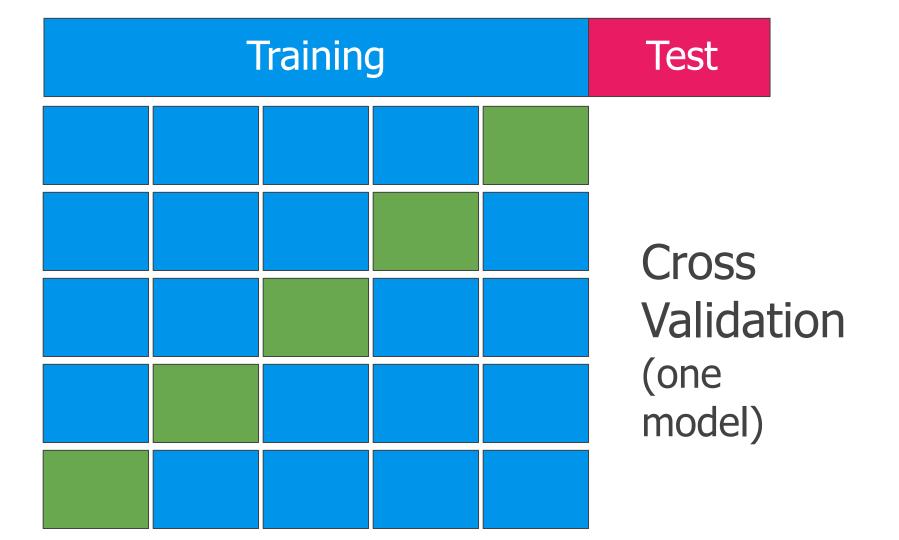
Pasting: sampling is performed without replacement.



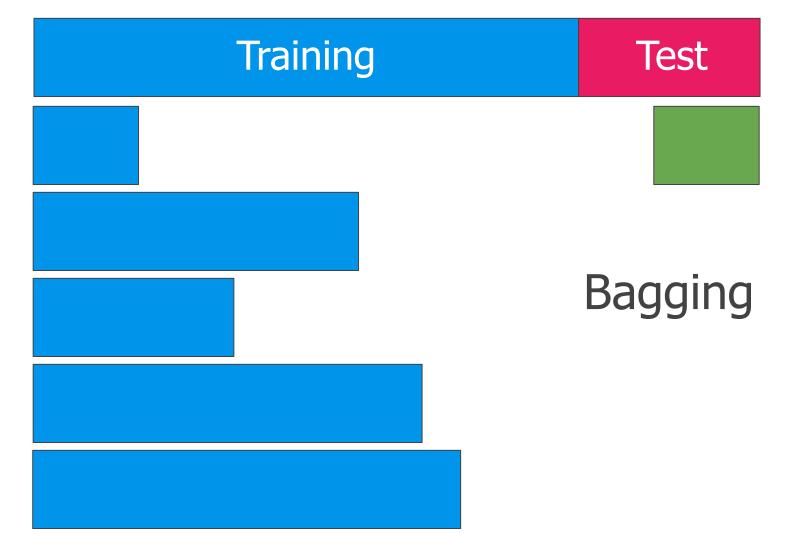
Cross Validation







Training Test Randomsubset Random subset Bagging Random subset (many models) Random subset Random subset



 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.

```
from sklearn.ensemble import BaggingClassifier
from sklearn.tree import DecisionTreeClassifier
bag clf = BaggingClassifier(
        DecisionTreeClassifier(),
        n estimators=500, max samples=100,
        bootstrap=True, n jobs=-1
bag clf.fit(X train, y train)
y pred = bag clf.predict(X test)
```

- Random Patches Ensemble method: sampling both training instances and features.
- This is particularly useful when dealing with high-dimensional inputs.

Ensemble Methods

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Boosting

 The general idea of most boosting methods is to train predictors sequentially, each trying to correct its predecessor.

Boosting

 The general idea of most boosting methods is to train predictors sequentially, each trying to correct its predecessor.

Most popular: AdaBoost and Gradient Boost.

AdaBoost [Freund and Schapire, 1997]

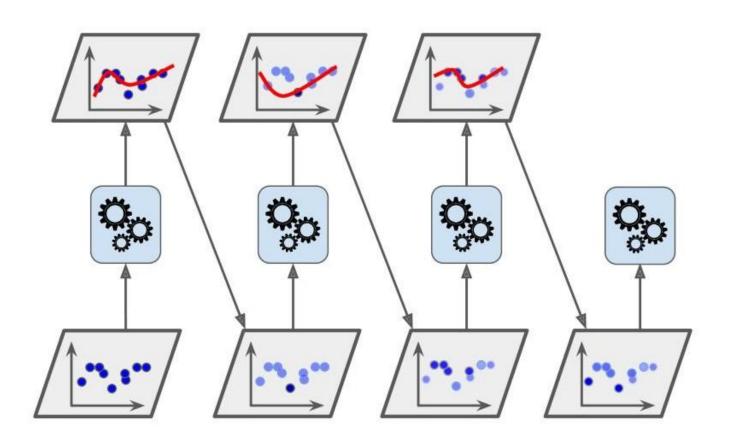
 One way for a new predictor to correct its predecessor is to pay a bit more attention to the training instances that the predecessor underfitted.

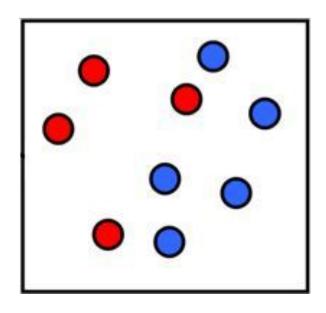
AdaBoost [Freund and Schapire, 1997]

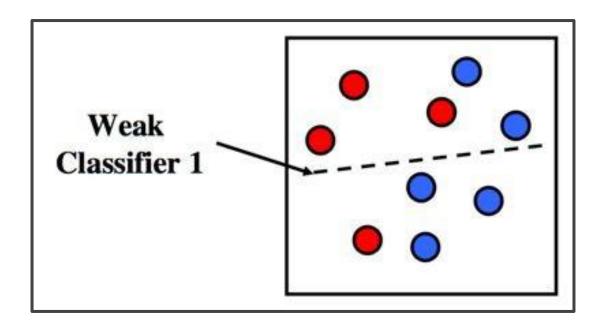
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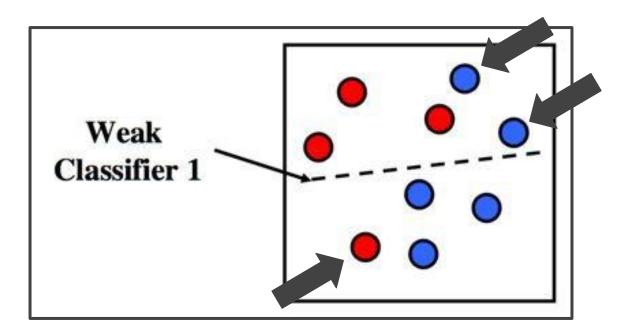
 This results in new predictors focusing more and more on the hard cases.

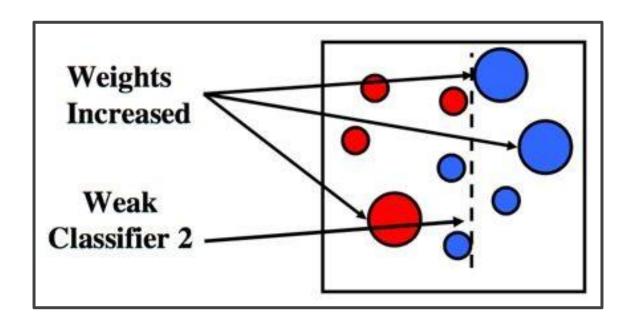
[&]quot;A Decision-Theoretic Generalization of On-Line Learning and an Application to Boosting", Y. Freund and R. Schapire (1997): http://goo.gl/OlduRW

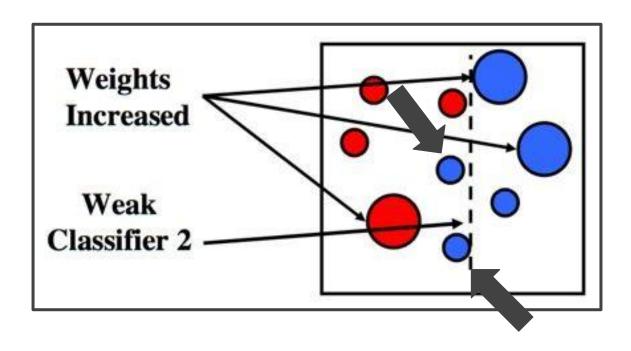


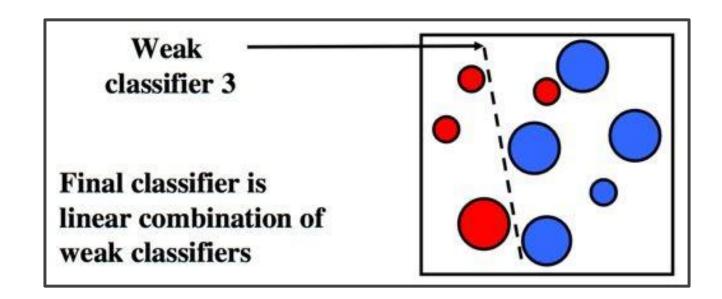


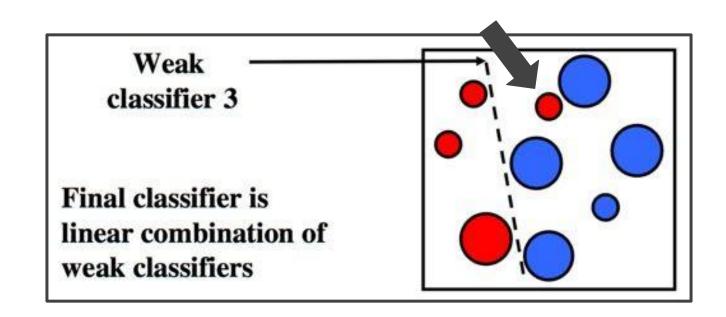


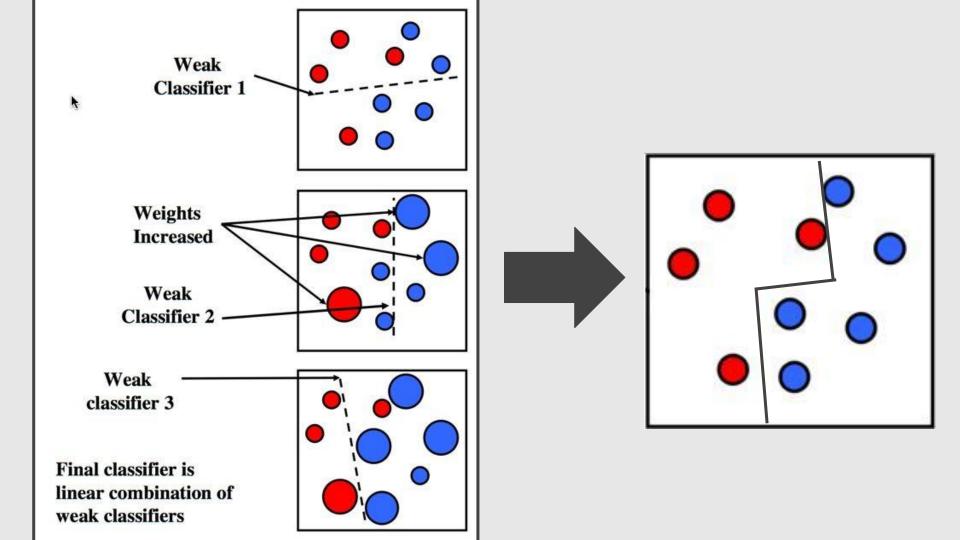










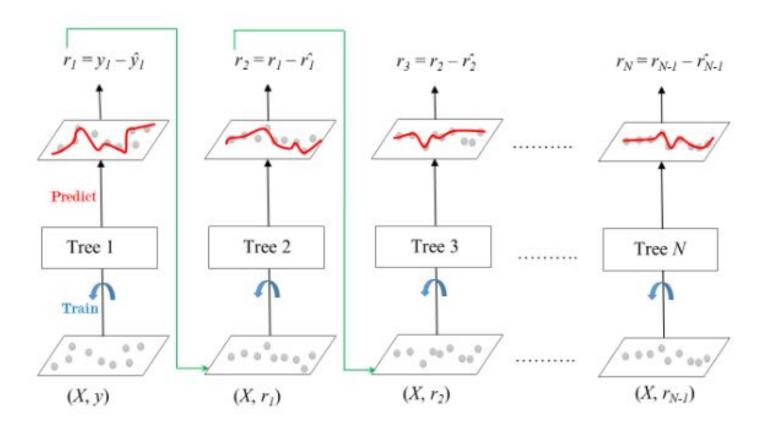




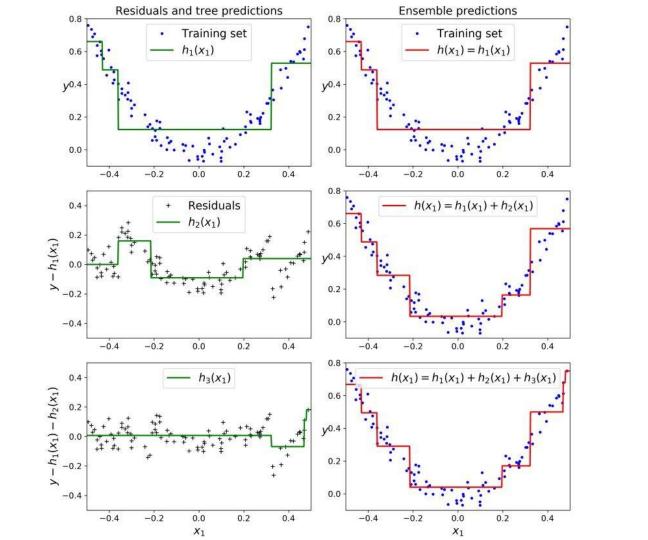
- 1. Assign every observation, x_i , an initial weight value, $w_i = \frac{1}{n}$, where n is the total number of observations.
- 2. Train a "weak" model. (most often a decision tree)
- 3. For each observation:
 - 3.1. If predicted incorrectly, wi is increased 3.2. If predicted correctly, wi is decreased
- 4. Train a new weak model where observations with greater weights are given more priority.
- 5. Repeat steps 3 and 9 until abservations perfectly predicted or a preset number of trees are trained.

 Instead of tweaking the instance weights at every iteration like AdaBoost does, this method fit the new predictor to the residual errors made by the previous predictor.

- Instead of tweaking the instance weights at every iteration like AdaBoost does, this method fit the new predictor to the residual errors made by the previous predictor.
- Instead of training on a newly sample distribution, the weak learner trains on the remaining errors.



$$y(pred) = y1 + (eta * r1) + (eta * r2) + + (eta * rN)$$



- 1. Fit a simple linear regressor or decision tree on data [call x as input and y as output]
- Calculate error residuals. Actual target value, minus predicted target value
 [e1 = y y_predicted1]
- 3. Fit a new model on error residuals as target variable with same input variables [call it e1 predicted]
- Add the predicted residuals to the previous predictions
 [y_predicted2 = y_predicted1 + e1_predicted]
- 5. Fit another model on residuals that is still left, i.e. **[e2 = y y_predicted2]** and repeat steps 2 to 5 until it starts overfitting or the sum of residuals become constant.

XGboost [Chen and Guestrin, 2016]:

Extreme Gradient Boosting

https://github.com/tgchen/xgboost

It aims at being extremely fast, scalable and portable.

Ensemble Methods

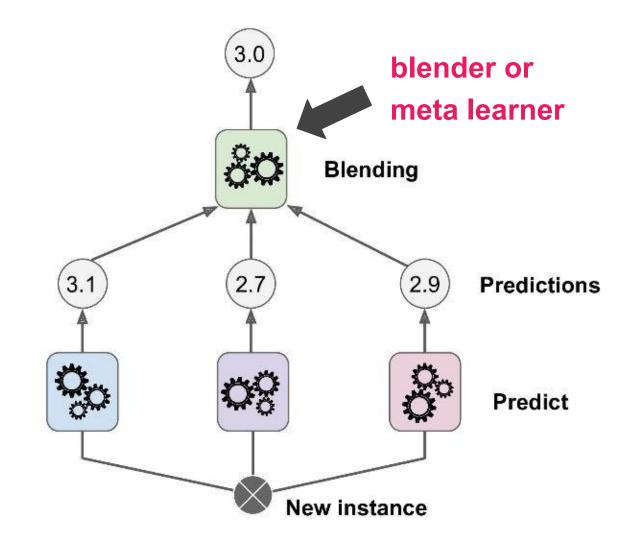
- Bagging (and Pasting)
- Boosting
- Stacking

Stacking [Wolpert, 1992]

Stacking (short for Stacked Generalization)

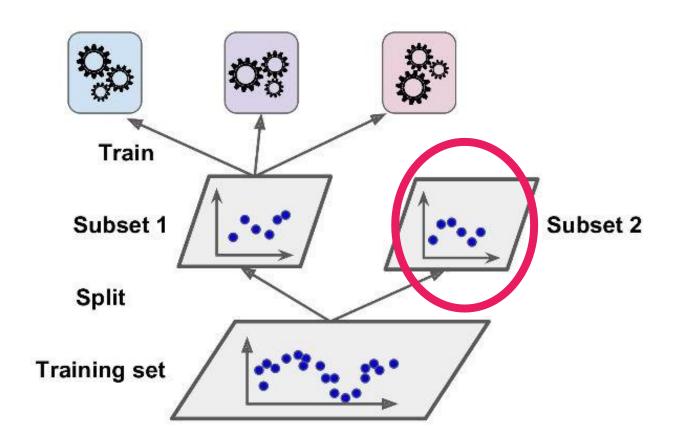
Stacking [Wolpert, 1992]

- Stacking (short for Stacked Generalization)
- Instead of using trivial functions (such as hard voting) to aggregate the predictions of all predictors in an ensemble, we train a model to perform this aggregation.

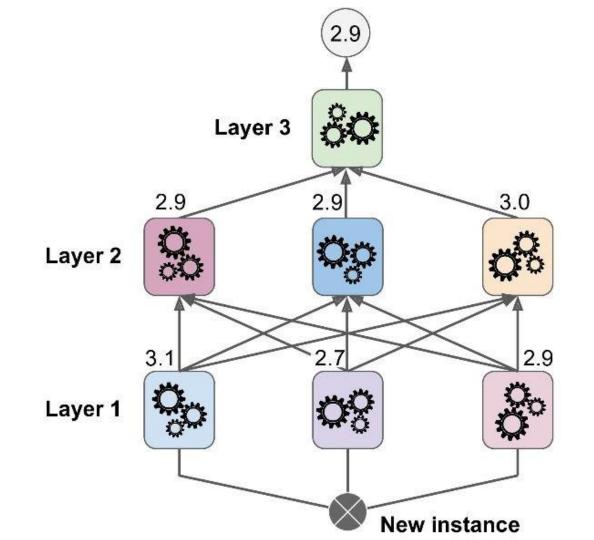


Stacking

To train the blender, a common approach is to use a hold-out set.



Multi-layer Stacking Ensemble



Stacking [Wolpert, 1992]

Scikit-Learn does not support stacking directly. =(

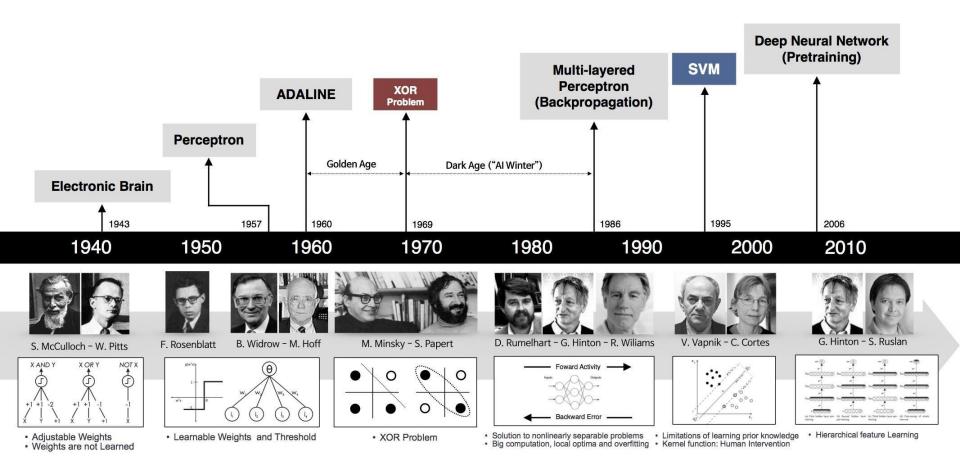
References

Machine Learning Books

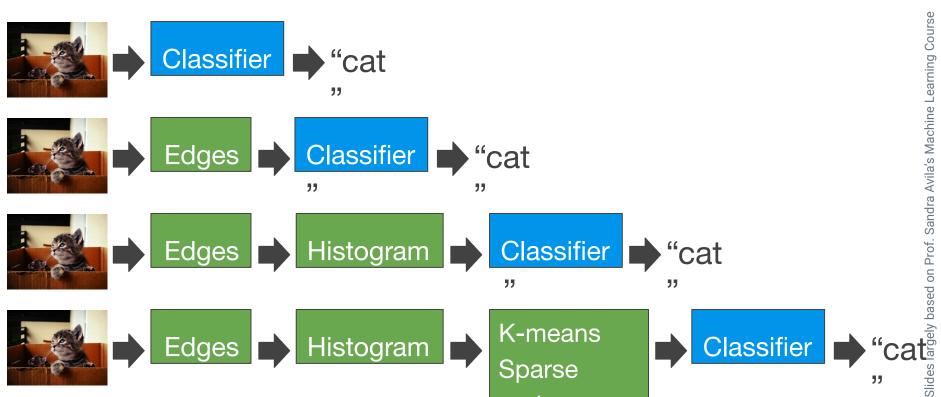
- Hands-On Machine Learning with Scikit-Learn and TensorFlow, Chap. 6 & 7
- Pattern Recognition and Machine Learning, Chap. 14
- Pattern Classification, Chap 8 & 9 (Sec. 9.5)
- "Scikit Learn Ensemble Learning, Bootstrap Aggregating (Bagging) and

Boosting" https://youtu.be/X3Wbfb4M33w

Support Vector Machines (SVMs)

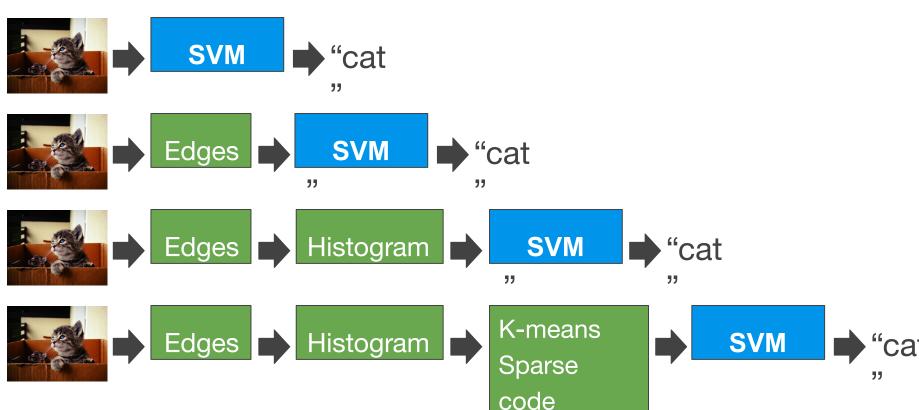


Traditional Recognition



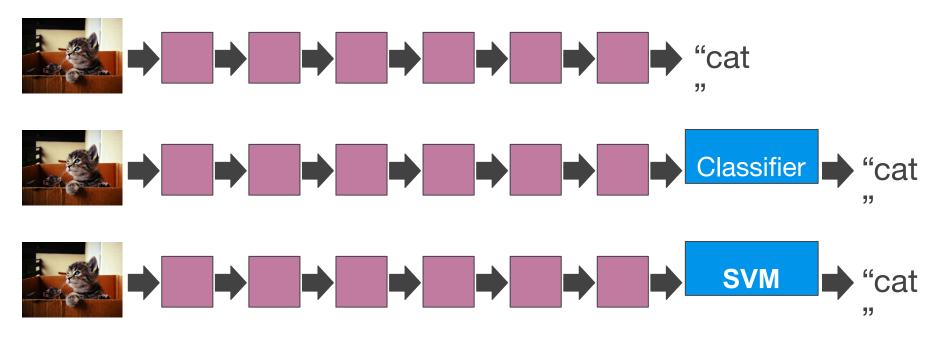
code

Traditional Recognition

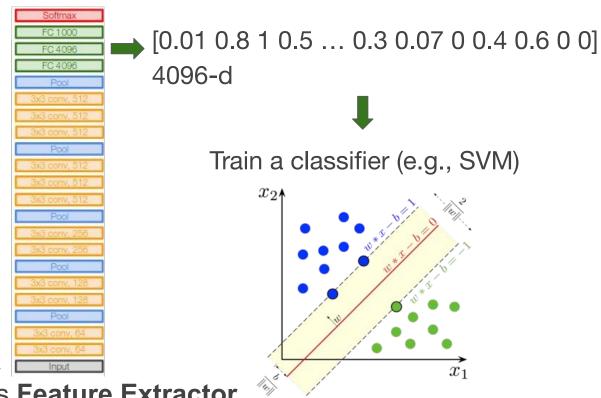


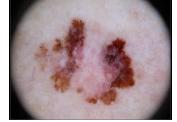
Slides largely based on Prof. Sandra Avila's Machine Learning Course

Deep Learning



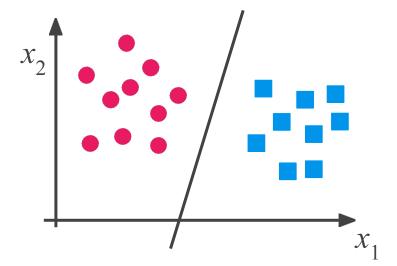
Transfer Learning with CNNs





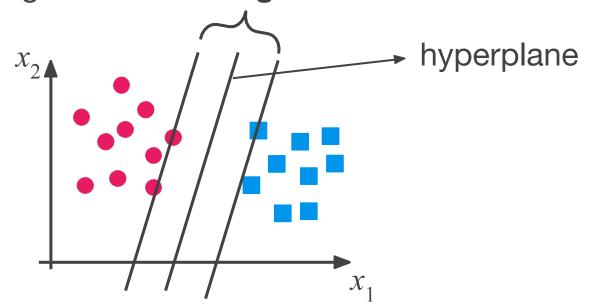
VGG as Feature Extractor

Idea of separating data with a large "gap".

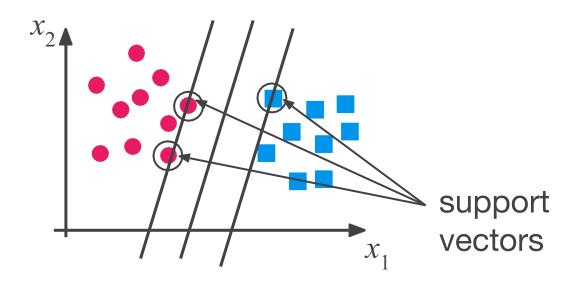


Idea of separating data with a large

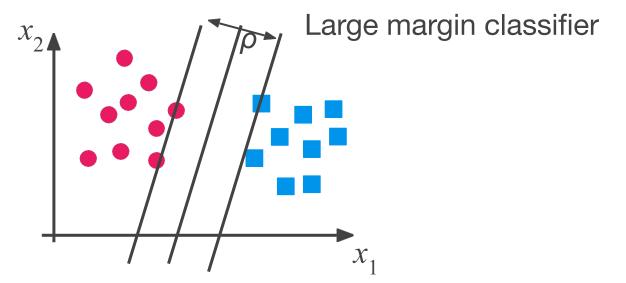
"gap".



Examples closest to the hyperplane are support vectors.

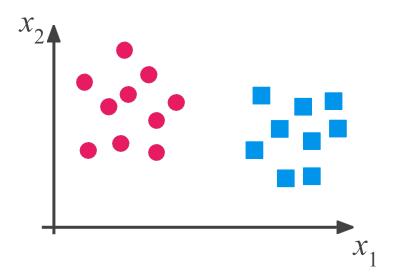


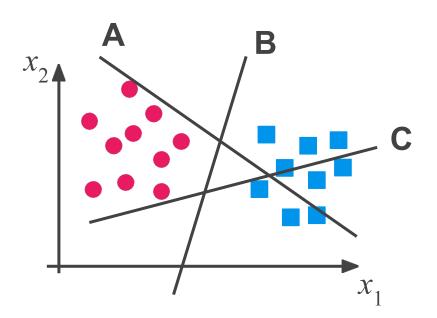
Margin ρ of the separator is the distance between support vectors.

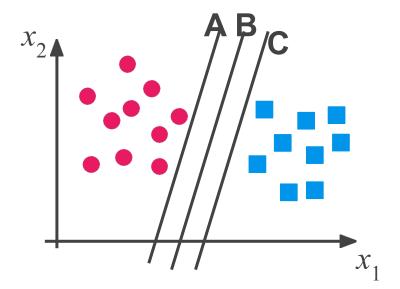


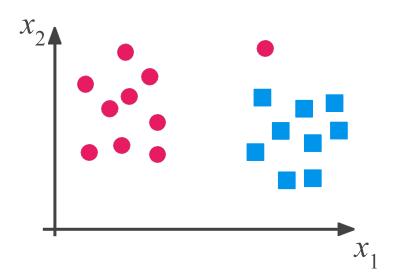
How does SVM work?

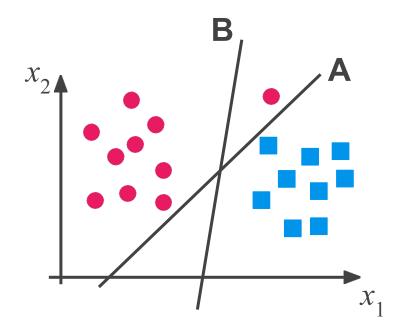
How can we identify the right hyperplane? Scenario 1

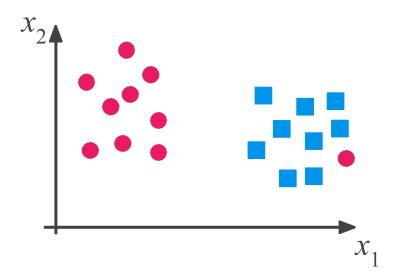


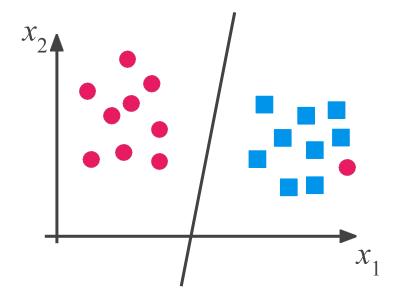






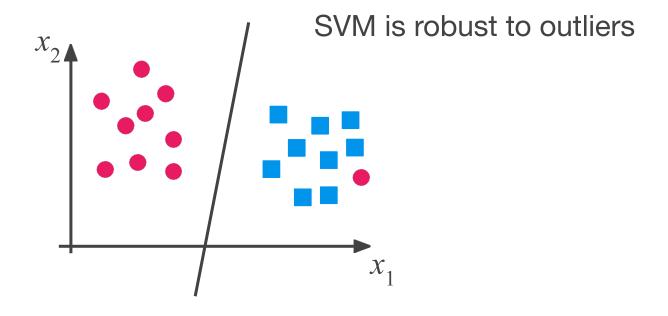


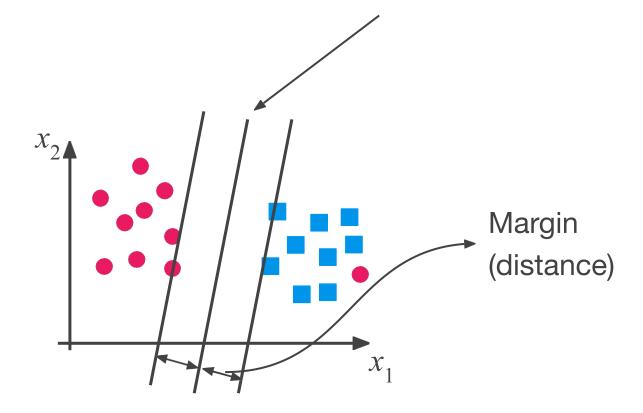




Scenario

4





SVM: Notation

We will be considering a linear classifier for a binary classification problem with labels y and features x.

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We will be considering a **linear classifier for a binary classification** problem with labels y and features x.

- Class labels: $y \in \{-1,1\}$ (instead of $\{0,1\}$)
- Parameters: w, b (instead of vector θ)

SVM: Notation

We will be considering a linear classifier for a binary classification problem with labels y and features x.

- Class labels: $y \in \{-1,1\}$ (instead of $\{0,1\}$)
- Parameters: w, b (instead of vector θ)
- Classifier: $h_{w,b}(x) = g(w^Tx + b)$
 - \circ g(z) = 1 if $z \ge 0$, and g(z) = -1 otherwise

Given a training example $(x^{(i)}, y^{(i)})$, we define the margin of (w, b) with respect to the training example:

$$y^{(i)}(w^Tx + b) \ge 1, i = \{1, ..., m\}.$$

Let $P(x^{(1)}, y^{(1)})$ be a point and l be a line defined by ax + by + c = 0. The distance d from P to l is defined by:

$$d(l,P) = \frac{|ax^{(1)} + by^{(1)} + c|}{\sqrt{a^2 + b^2}}$$

Let $P(x^{(1)}, y^{(1)})$ be a point and l be a line defined by ax + by + c = 0. The distance d from P to l is defined by:

$$d(l,P) = \frac{|ax^{(1)} + by^{(1)} + c|}{\sqrt{a^2 + b^2}}$$

$$d(w,b,x) = \frac{|w^Tx + b|}{||w||}$$

$$d(w,b,x) = |w^T x + b|$$

$$|w||$$



$$\min_{w,b} \frac{1}{2} ||w||^{2}$$
s.t. $y^{(i)}(w^{T}x + b) \ge 1, i = \{1, ..., m\}$

$$d(w,b,x) = |w^T x + b|$$

$$|w|$$

http://cs229.stanford.edu/notes2019fall/cs229-notes3.pdf

$$\min_{w,b} ||w||$$
s.t. $y^{(i)}(w^Tx + b) \ge 1, i = \{1, ..., m\}$

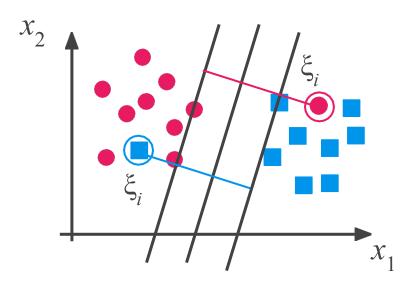
Need to optimize a quadratic function subject to linear

Soft Margin Classification

What if the training set is not linearly separable?

Soft Margin Classification

Slack variables ξ_i can be added to allow misclassification of difficult or noisy examples, resulting margin called **soft**.

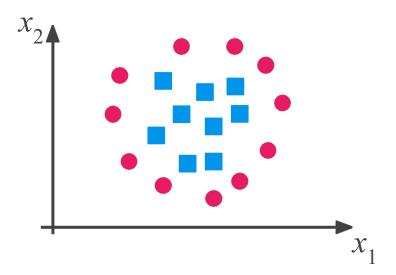


Soft Margin Classification

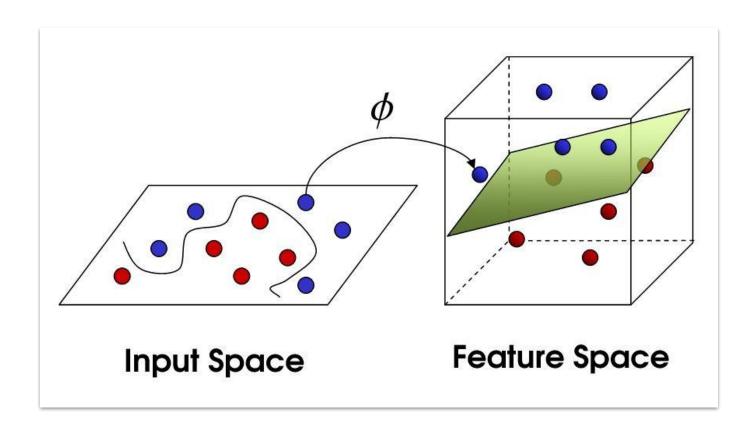
Modified formulation incorporates slack variables:

$$\min_{\substack{w,b,\xi\\s.t.\ y_i(w^Tx + b^n) \ge 1 - \xi_i,\ \xi_i \ge 0,\ i = \{1, ..., m\}}} \frac{1/2||w||^2 + C\sum_i \xi_i}{s.t.\ y_i(w^Tx + b^n) \ge 1 - \xi_i,\ \xi_i \ge 0,\ i = \{1, ..., m\}$$

Parameter *C* can be viewed as a way to control overfitting: it "trades off" the relative importance of maximizing the margin and fitting the training data.

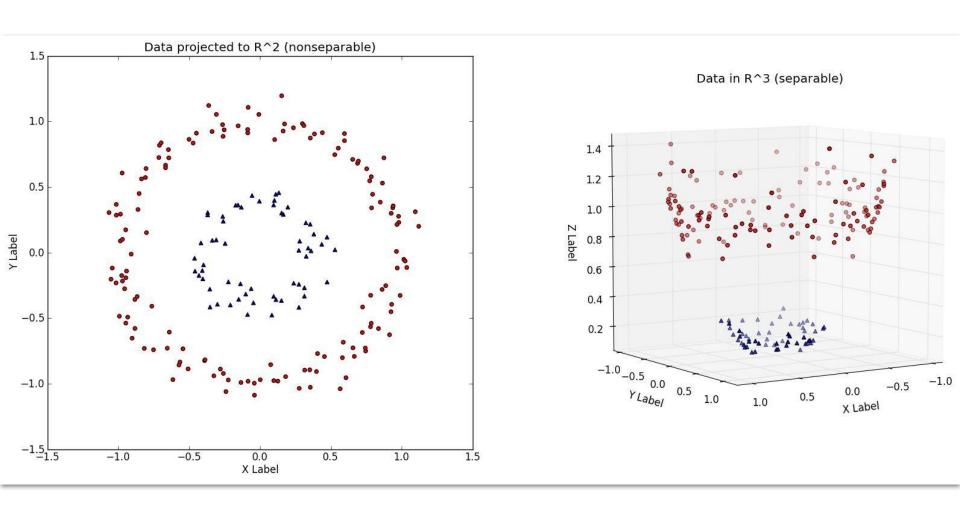


Kernel Trick



Kernel Trick

- Linear SVM: $x_i \cdot x_j$
- Nonlinear SVM: $K(x_i, x_j) = \phi(x_i) \cdot \phi(x_i)$, feature mapping ϕ
- Kernel matrix $K_{ij} = K(x_i, x_j) = \phi(x_i) \cdot \phi(x_j) = \phi(x_j) \cdot \phi(x_i) = K_{ji}$
- Radial Basis Function (RBF) kernel
- Gaussian kernel
- Polynomial kernel
- Chi-square kernel, histogram intersection kernel, string kernel,



Important Parameters

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

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C: Penalty parameter C of the error term. It also controls the trade off between smooth decision boundary and classifying the training points correctly.

Important Parameters

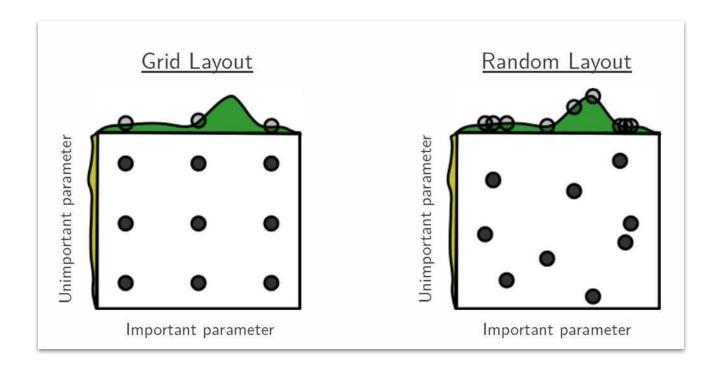
Important parameters having higher impact on model performance, "kernel", "gamma" and "C".

C: Penalty parameter C of the error term. It also controls the trade off between smooth decision boundary and classifying the training points correctly.

The parameters can be tuned using grid-search.



Grid Search



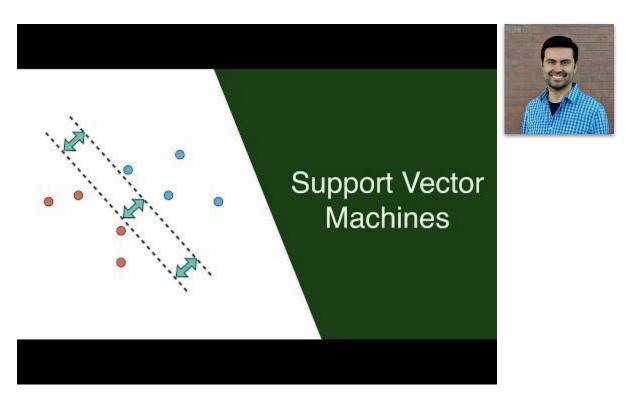
[&]quot;Random Search for Hyper-Parameter Optimization": http://www.jmlr.org/papers/volume13/bergstra12a/bergstra12a.pdf

Libraries

- Scikit-learn: https://scikit-learn.org/stable/modules/svm.html
- LIBSVM: https://www.csie.ntu.edu.tw/~cjlin/libsvm
- LIBLINEAR: https://www.csie.ntu.edu.tw/~cjlin/liblinear
- PmSVM:

https://sites.google.com/site/wujx2001/home/power-mean-svm

Support Vector Machines (SVMs): A friendly introduction



https://www.youtube.com/watch?v=Lpr X8zuE8

References

Machine Learning Books

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

Machine Learning Courses

- https://www.coursera.org/learn/machine-learning, Week 7
- http://cs229.stanford.edu/notes2019fall/cs229-notes3.pdf

Decision Trees

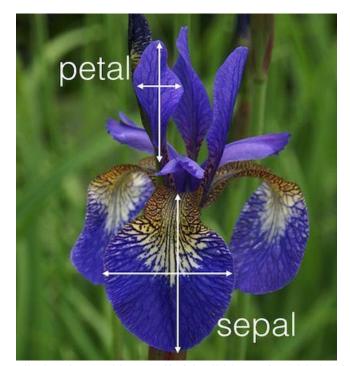
Decision Tree & Random Forest

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

Decision Tree & Random Forest

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

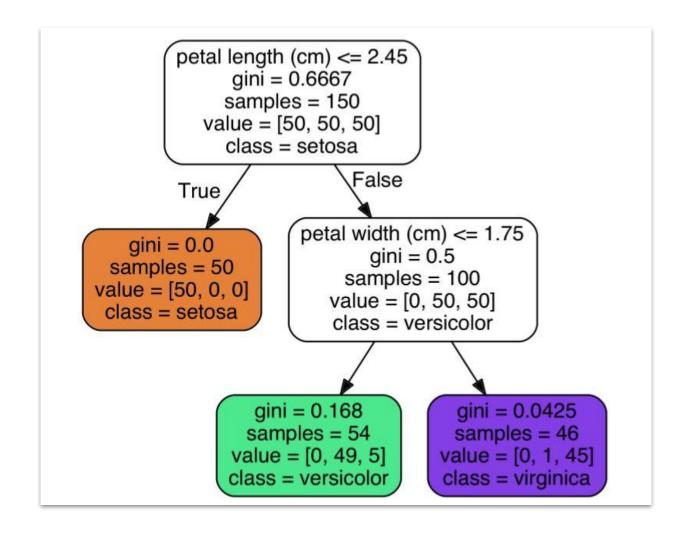
150 iris flowers from three different species.

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:

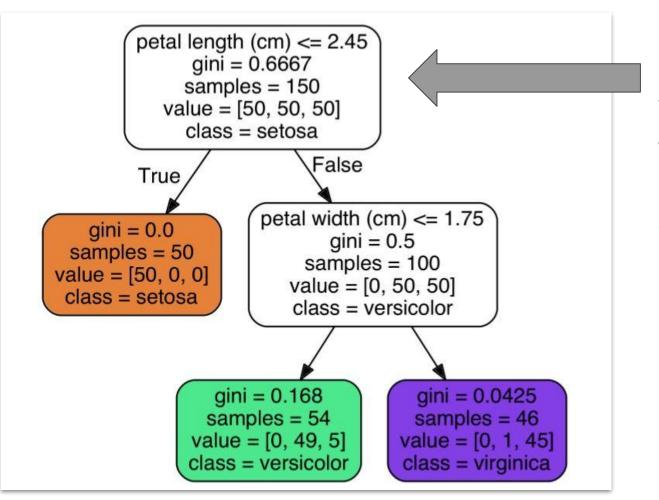
- sepal length in cm
- 2. sepal width in cm
- B. petal length in cm
- 4. petal width in cm



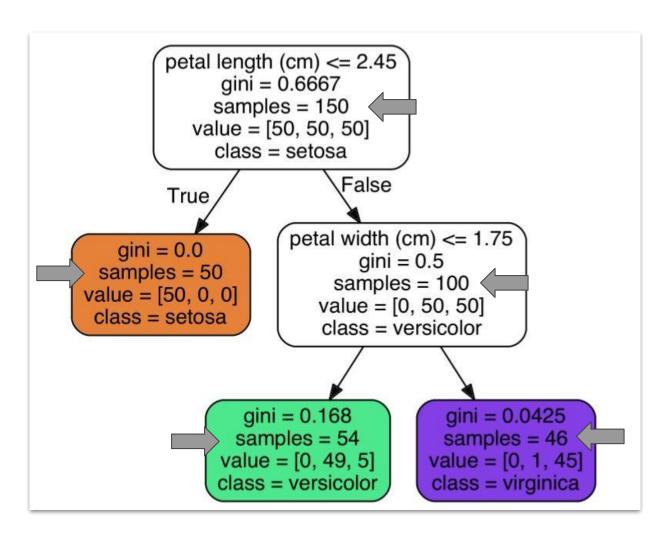
```
iris = load iris()
X = iris.data[:, 2:] # petal length and width
y = iris.target
tree clf = DecisionTreeClassifier(max depth ≥)
tree clf.fit(X, y)
#You can visualize the trained Decision Tree by first using the export graphviz() method
to output a graph definition file called iris tree.dot:
from sklearn.tree import export graphviz
export graphviz(
    tree clf,
    out file="iris tree.dot",
    feature names=iris.feature names[2:],
    class names=iris.target names,
    rounded=True,
    filled=True
$ dot -Tpng iris tree.dot -o iris tree.png#convert this .dot file to a variety of
formats such as PDF or PNG using the dot command-line tool from the graphviz package
```

from sklearn.datasets import load iris

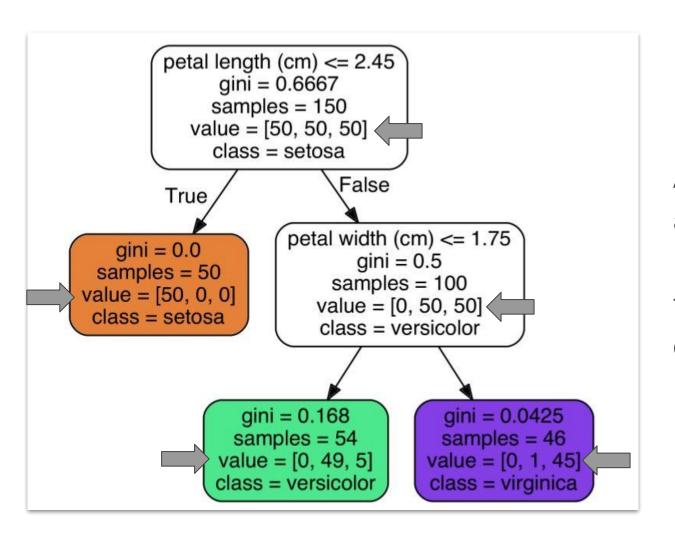
from sklearn.tree import DecisionTreeClassifier



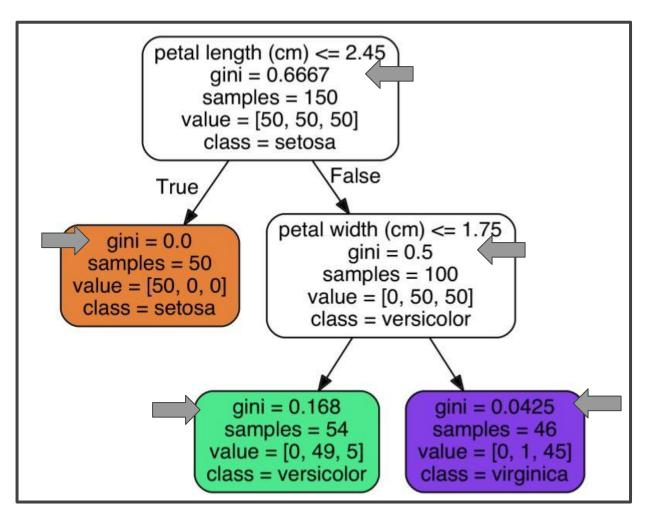
This node asks whether the flower's petal length is smaller than 2.45 cm



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.



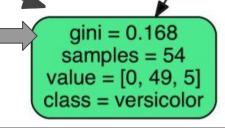
A **node's gini** attribute measures its impurity.

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

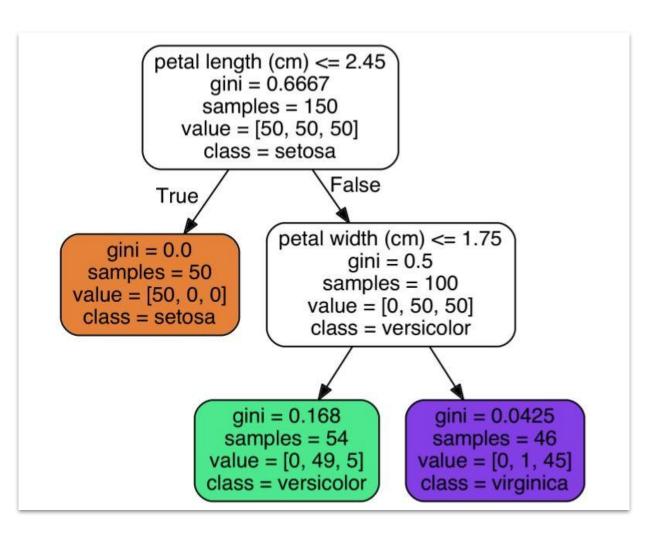
For example, the depth 2 left node has a gini score equal to $1 - (0/54)^2 - (49/54)^2 - (5/54)^2 \approx 0.168$.

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

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

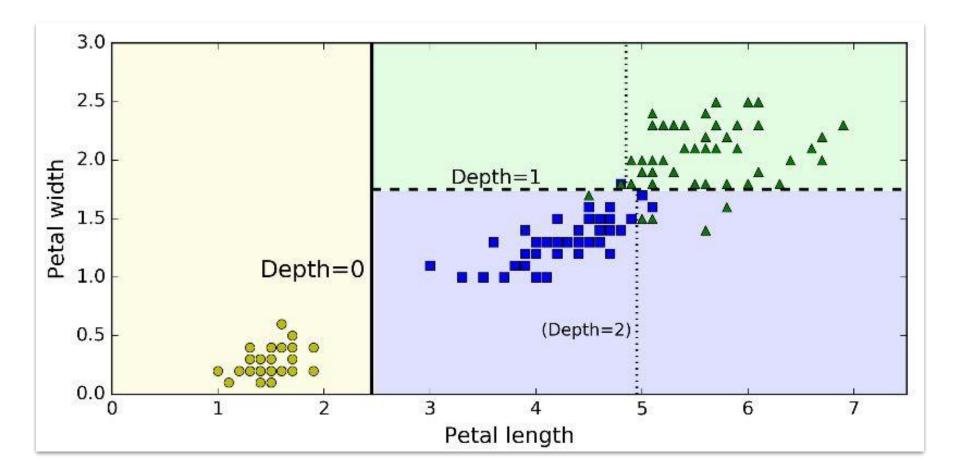


gini = 0.0425 samples = 46 value = [0, 1, 45] class = virginica belong to the same class.



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

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



Model Interpretation

- White Box vs Black Box
- Decision Trees are fairly intuitive and their decisions are easy to interpret (white box models).
- Random Forests or neural networks are generally considered black box models.

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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- How does it choose k and t_k ?

- Classification And Regression Tree (CART) algorithm.
- 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 ?

 It searches for the pair (k, t_k) that produces the purest subsets (weighted by their size).

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

CART cost function for classification

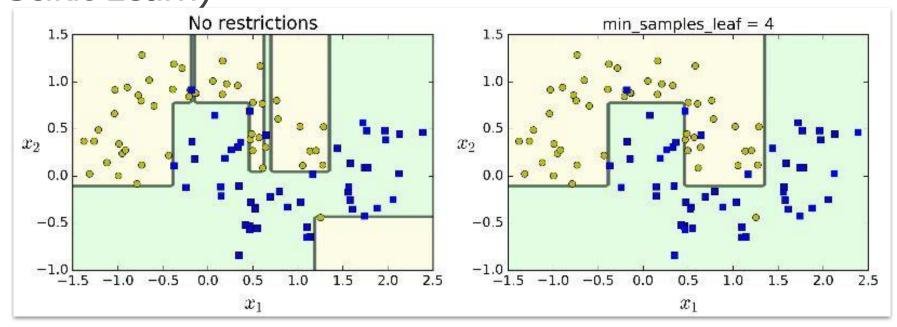
It stops recursing once it reaches the maximum depth (hyperparameter), or if it cannot find a split that will reduce impurity

- CART algorithm is a greedy algorithm.
- Training: O(n × m log(m)), m samples, n features.
- Overall prediction: O(log2 (m)).

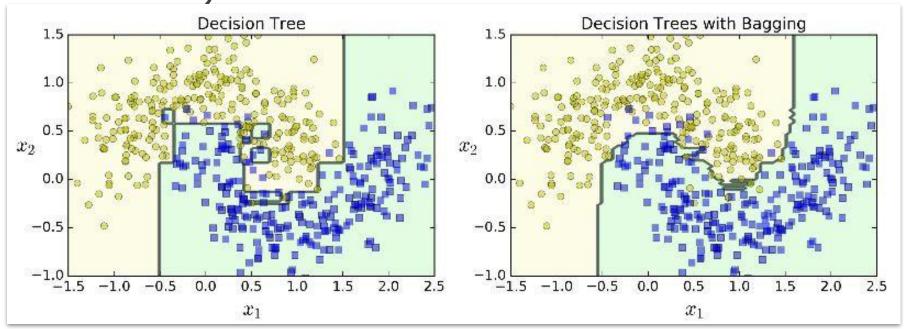
Regularization (in Scikit-Learn)

- max_depth hyperparameter: the default value is None,
 which means unlimited.
- min_samples_split: the minimum number of samples a node must have before it can be split.
- min_samples_leaf: the minimum number of samples a leaf node must have.

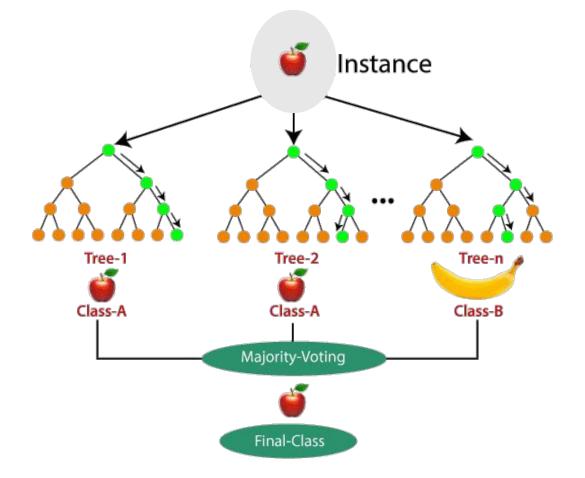
Regularization (in Scikit-Learn)



Regularization (in Scikit-Learn)



Random Forest



 Random Forest is an ensemble of Decision Trees, generally trained using the Bagging method.

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• Extra randomness when growing trees:

 Instead of searching for the very best feature when splitting a node, it searches for the best feature among a random subset of features.

1. Assume number of cases in the training set is *N*. Then, sample of these *N* cases is taken at random but with replacement.

2. If there are *M* input variables, a number *m*<*M* is specified such that at each node, *m* variables are selected at random out of the *M*.

The best split on these m is used to split the node. The value of m is held constant while we grow the forest.

- Each tree is grown to the largest extent possible and there is no pruning.
- 4. Predict new data by aggregating the predictions of the ntree trees (i.e., majority votes for classification, average for regression).

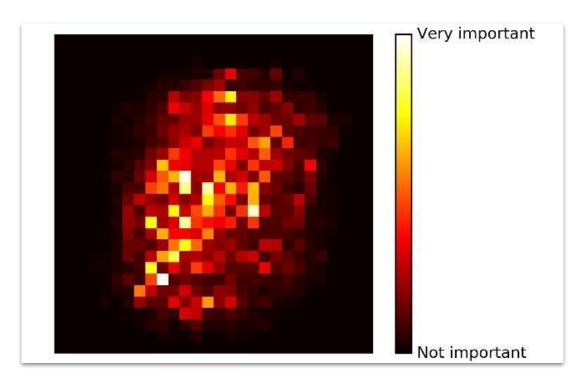
Random Forest: Feature Importance

(cm) 0.423357996355

```
from sklearn.datasets import load_iris
from sklearn.ensemble import RandomForestClassifier iris =
load_iris()
rnd_clf = RandomForestClassifier(n_estimators= 500, n_jobs= -1)
rnd_clf.fit(iris[ "data"], iris["target"])
for name, score in zip(iris["feature_names"],
rnd_clf.feature_importances_):
    print(name, score)
```

```
sepal length (cm)
0.112492250999 sepal width (cm)
0.0231192882825 petal length
(cm) 0.441030464364 petal width
```

Random Forest: Feature Importance



MNIST pixel importance (according to a Random Forest classifier)

Forward Thinking: Building Deep Random Forests

Kevin Miller, Chris Hettinger, Jeffrey Humpherys, Tyler Jarvis, and David F

Department of Mathematics Brigham Young University Provo, Utah 84602

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Abstract

Training Big Random Forests with Little Resources

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Department of Computer Science
University of Copenhagen
Copenhagen, Denmark
fabian.gieseke@di.ku.dk

ABSTRACT

Without access to large compute clusters, building random forests on large datasets is still a challenging problem. This is, in particular, the case if fully-grown trees are desired. We propose a simple yet effective framework that allows to efficiently construct ensembles of huge trees for hundreds of millions or even billions of training instances using a cheap desktop computer with commodity hardware. The basic idea is to consider a multi-level construction scheme, which builds top trees for small random subsets of the available data and which subsequently distributes all training instances to the top trees' leaves for further processing. While being conceptually simple, the overall efficiency crucially depends on the particular implementation of the different phases. The practical merits of our

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ensembles in a parallel or distridividual compute nodes (e.g., by node). While this can significat such frameworks naturally requing environments. Further, the el might cause problems in case the large to fit into the main memoi

In this work, we propose a scheme for building random for scale. The main idea is to build phases: Starting with a top tree be the data, one subsequently distresses of that tree. For each leaf

Distributed Deep Forest and its Application to Automatic Detection of Cash-out Fraud

Ya-Lin Zhang[†], Jun Zhou[‡], Wenhao Zheng[†], Ji Feng[†], Longfei Li[‡], Ziqi Liu[‡], Ming Li[†], Zhiqiang Zhang[‡], Chaochao Chen[‡], Xiaolong Li[‡], Zhi-Hua Zhou[†]

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‡Ant Financial Services Group, China

‡{jun.zhoujun, longyao.llf, ziqiliu, lingyao.zzq, chaochao.ccc, xl.li}@antfin.com

Deep Forest: Towards an Alternative to Deep Neural Networks*

Zhi-Hua Zhou and Ji Feng

National Key Lab for Novel Software Technology, Nanjing University, Nanjing 210023, China {zhouzh, fengj}@lamda.nju.edu.cn

Abstract

2018

May

In this paper, we propose gcForest, a decision tree

ample, even when several authors all use convolutional neural networks [LeCun *et al.*, 1998; Krizhenvsky *et al.*, 2012; Simonyan and Zisserman, 2014], they are actually using dif-

References

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)

https://towardsdatascience.com/random-forest-in-python-24d0893d51c0