

CAS Practical Machine Learning Introduction

**Project: First ML Steps with Python** 

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# scikit-learn

Machine Learning with Python

## Machine Learning with scikit-learn

scikit-learn (http://scikit-learn.org/stable/index.html)

- provides many ML algorithms
- plus data preprocessing, feature selection, evaluation, visualization
- open source with BSD license
- installation
  - pre-installed with Anaconda
  - or see https://scikit-learn.org/stable/install.html
- general documentation see <a href="http://scikit-learn.org/stable/documentation.html">http://scikit-learn.org/stable/documentation.html</a>

# Basic Workflow with scikit-learn (1)

- 1) loading data (http://scikit-learn.org/stable/datasets/index.html)
  - dataset ds needs to have a specific format
    - ds.data is an array [n\_samples, n\_features] for storing the relevant features (numeric)
    - for supervised training: ds.target is an array [n\_samples] with preassigned classes (numeric)
  - prepared random ("generated"), small ("toy") and large ("real-world")
    datasets exist
- 2) training the model (<a href="http://scikit-learn.org/stable/user\_guide.html">http://scikit-learn.org/stable/user\_guide.html</a>)
  - each supervised ML algorithm ml ("estimator") implements the method fit (ds.data, ds.target) that takes a training dataset ds
- 3) using the model (<a href="http://scikit-learn.org/stable/user\_guide.html">http://scikit-learn.org/stable/user\_guide.html</a>)
  - each ML algorithm ml implements the method predict (ds.data) that takes an array ds.data and returns the predicted classes as array [n samples]

# Basic Workflow with scikit-learn (2)

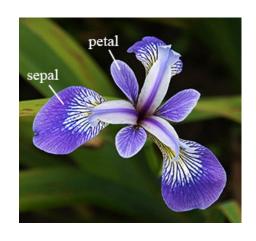
- 4) evaluating the model (<a href="http://scikit-learn.org/stable/modules/model\_evaluation.html">http://scikit-learn.org/stable/modules/model\_evaluation.html</a>)
  - each ML algorithm ml has a method score (ds.data, ds.target) to calculate a model-specific, commonly-used evaluation metric
  - more elaborate evaluation tools support different metrics
- 5) saving and loading the trained model
  - via joblib (https://pythonhosted.org/joblib/persistence.html)
  - joblib.dump(ml, filename) and joblib.load(filename)

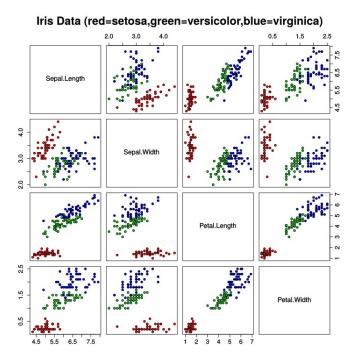
### **Decision Tree**

Baseline Supervised Classification Algorithm

#### Iris Flower Dataset

- samples of different species of iris flowers
  - 3 species: Iris setosa, Iris virginica and Iris versicolor)
  - 4 features: sepal/petal length/width in centimeters
- introduced in the article: R. A. Fisher (1936).
   "The use of multiple measurements in taxonomic problems". Annals of Eugenics. 7 (2): 179-188
- see <u>https://en.wikipedia.org/wiki/Iris\_flower\_dat</u> a\_set
- dataset is predefined in scikit-learn
   (<a href="http://scikit-learn.org/stable/modules/generated/sklearn.datasets.load\_iris.html">http://scikit-learn.datasets.load\_iris.html</a>)





### Decision Tree with scikit-learn (1)

#### Jupyter Notebook

1) loading the dataset

```
from sklearn import datasets
iris = datasets.load iris()
```

- 2) exploring and visualizing the dataset
  - raw dataset

```
print(iris.data)
```

features

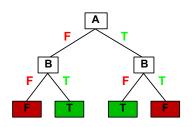
```
print(iris.feature names)
```

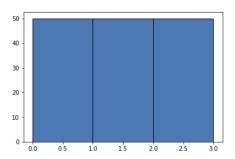
classes

```
print(iris.target)
print(iris.target.size)
print(iris.target names)
```

classes histogram

```
import matplotlib.pyplot as plt
plt.hist(iris.target, bins=(0,1,2,3),
        edgecolor='black')
plt.show()
```





### Decision Tree with scikit-learn (2)

- 2) exploring and visualizing the dataset
  - scatterplot with sepal length and width

```
import matplotlib.pyplot as plt
X = iris.data[:, :2] # only the first two features
y = iris.target
plt.figure(2, figsize=(8, 6))
plt.clf()
plt.scatter(X[:, 0], X[:, 1], c=y, cmap=plt.cm.Set1,
edgecolor='k')
plt.xlabel('Sepal length')
plt.ylabel('Sepal width')
plt.xticks(())
plt.yticks(())
plt.show()
```

### Decision Tree with scikit-learn (3)

- 3) training the model (<a href="http://scikit-learn.org/stable/modules/tree.html">http://scikit-learn.org/stable/modules/tree.html</a>)
  - scikit-learn implements the CART DT algorithm
    - recursively splits the dataset into subsets based on conditions wrt feature values
    - minimizes the Gini impurity to decide on the best split
      - measures the error rate that would result from classifying the remaining data samples randomly according to the distribution of labels in the candidate subset
      - i.e., the error rate that would result if the tree does not grow any further
  - running the classifier

```
from sklearn.tree import DecisionTreeClassifier
classifier = DecisionTreeClassifier()
# use all but the last sample for training
classifier.fit(iris.data[:-1], iris.target[:-1])
```

4) evaluating the model

```
classifier.predict(iris.data[-1:])
```

### Decision Tree with scikit-learn (4)

- 5) visualizing the model (optional)
  - for rendering, we use Graphviz (<a href="http://www.graphviz.org/">http://www.graphviz.org/</a>)
    - general tool for visualizing graphs and networks
    - download from <a href="http://www.graphviz.org/download/">http://www.graphviz.org/download/</a>
    - install and verify via dot -V
  - option 1: direct display

- option 2: render image file via command line
  - in Python, save tree in Graphviz file format ("tree.dot")

then render file via command line

```
dot -Tpng tree.dot -o tree.png
```

