PATTERN RECOGNITION USING PYTHON

Classification

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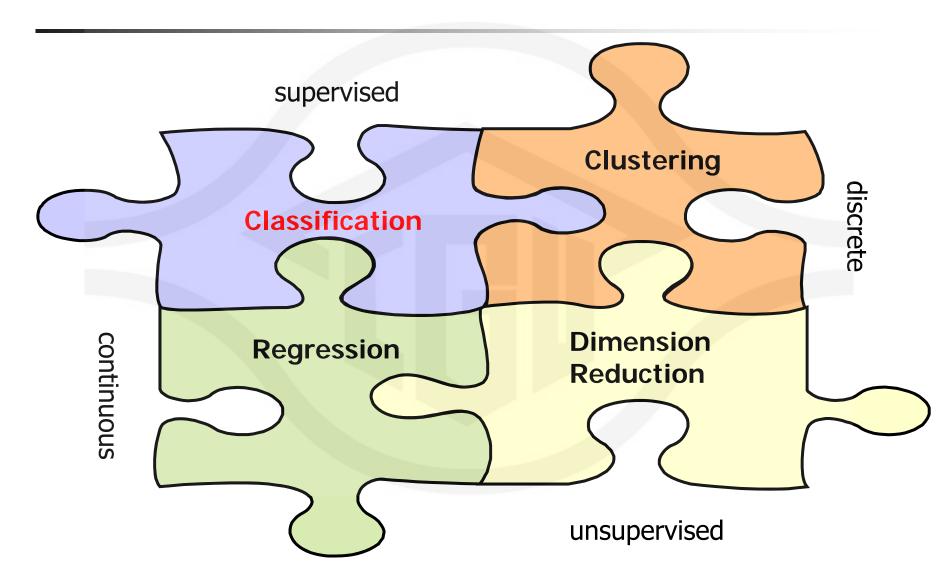
Chang Gung University, Taiwan

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Supervised Learning Algorithms for Classification

- Popular algorithms for classification, such as logistic regression, support vector machines, and decision trees
- Examples and explanations using the scikit-learn machine learning library, which provides a wide variety of machine learning algorithms via a user-friendly Python API

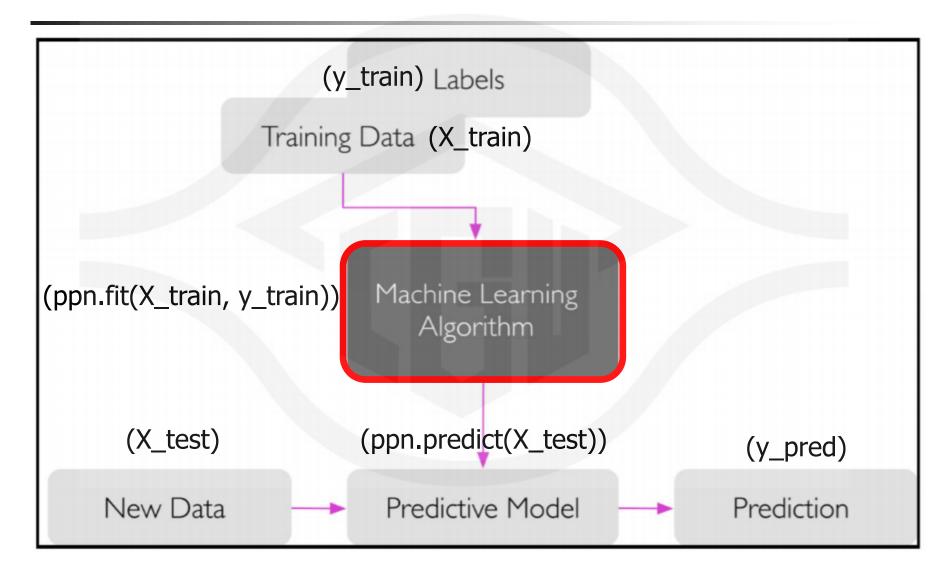
Machine Learning Organizational Chart



Choosing a Classification Algorithm

- Selecting features and collecting training samples
- Choosing a performance metric
- Choosing a classifier and optimization algorithm
- Evaluating the performance of the model
- Tuning the algorithm

Solve Classification Problem



Problem Transformation

Machine Learning Algorithm

Classification Problem



evaluate **error** (when label ≠ prediction) by **cost function** (objective function)

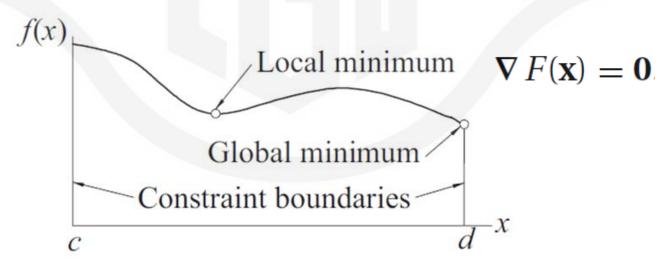
Optimization Problem

Optimization Problem

- In engineering, optimization is closely related to design.
- We wish to keep it (objective function) as small as possible, such as the cost or weight.

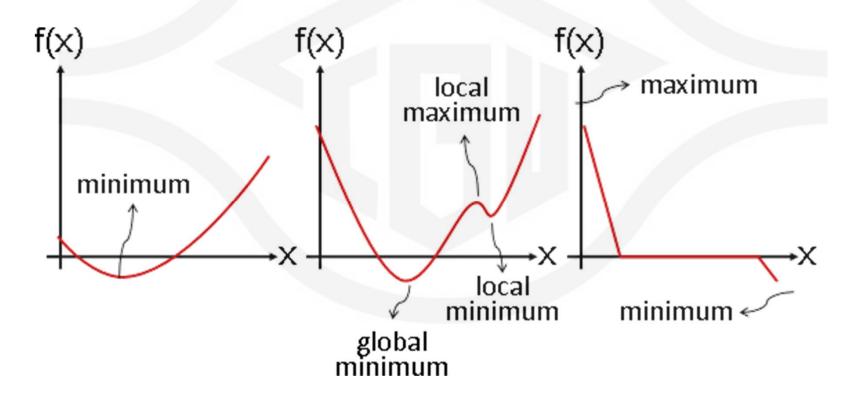
Find **x** that minimizes
$$F(\mathbf{x})$$
 subject to $g(\mathbf{x}) = 0$, $h(\mathbf{x}) \ge 0$.

Find the points where the gradient vector of F(x) vanishes

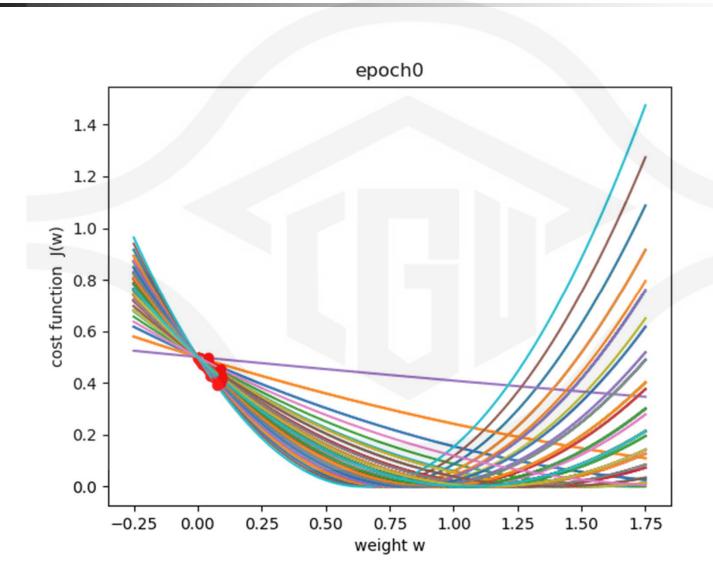


Algorithms for Minimization

- It require starting values (self.w_)
- Processes by iterative procedures (n_iter)

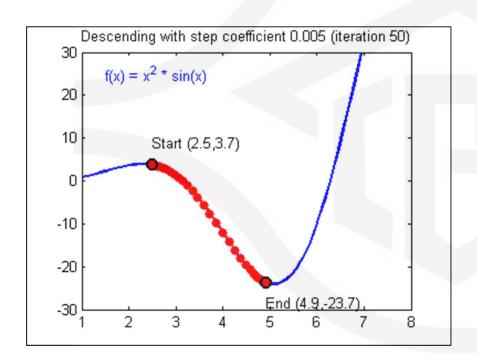


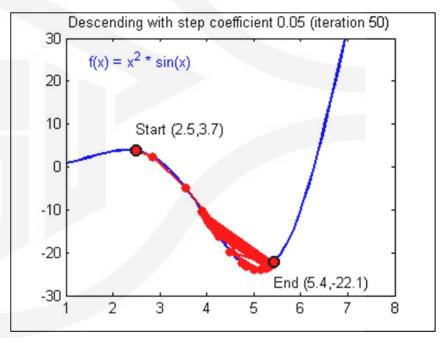
Gradient Descent to Achieve Minimization



Effect of Learning Rate

Gradient descent with different learning rate



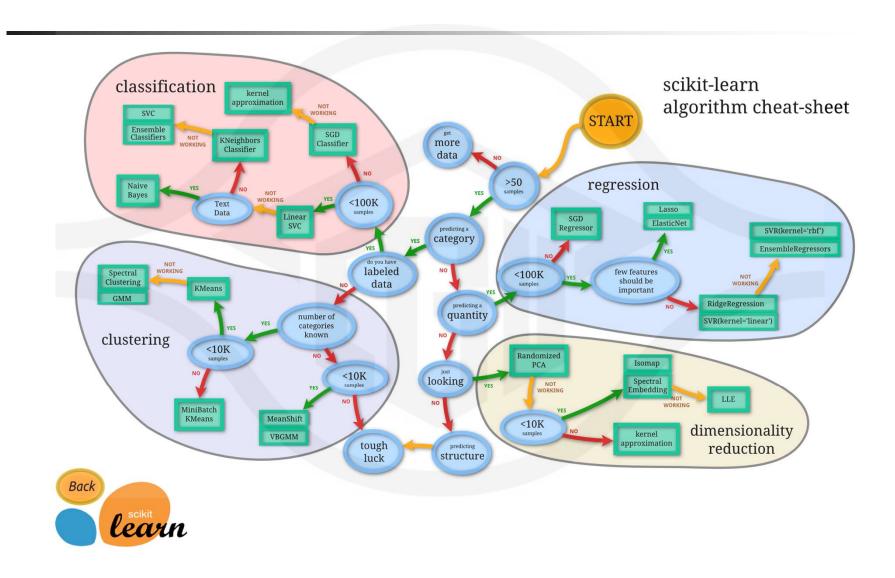


What is Scikit-Learn

https://scikit-learn.org/stable/



Learning Map of Scikit-Learn



Import Scikit-Learn (sklearn)

from ... import ... (special use method)

```
from sklearn import datasets
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
from sklearn.linear_model import Perceptron
from sklearn.metrics import accuracy_score
```

Load Data via sklearn

Load data by built-in scikit-learn datasets

```
iris = datasets.load_iris()
X = iris.data[:, [2, 3]]
y = iris.target
print('Class labels:', np.unique(y))
```

The iris dataset is a classic and very easy multi-class classification dataset.

Classes	3
Samples per class	50
Samples total	150
Dimensionality	4
Features	real, positive

Preprocessing via sklearn

Split data (70% train set and 30% test set)

```
X_train, X_test, y_train, y_test = train_test_split(
    X, y, test_size=0.3, random_state=1, stratify=y)
```

Standardize features

```
sc = StandardScaler()
sc.fit(X_train)
X_train_std = sc.transform(X_train)
X_test_std = sc.transform(X_test)
```

Training and Testing

Initialize the object and training

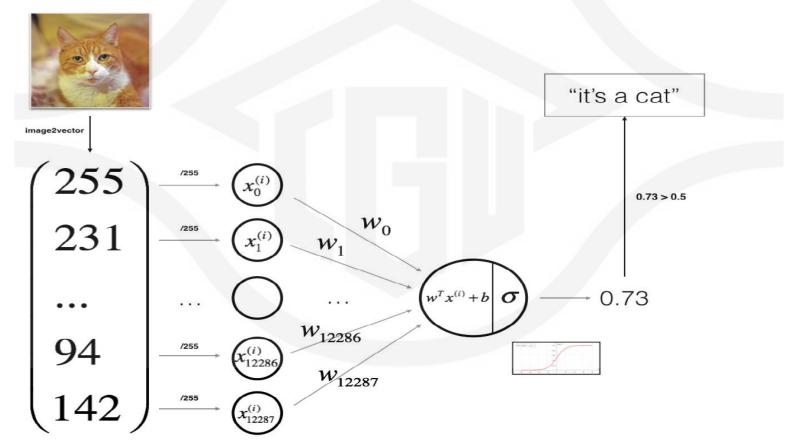
```
ppn = Perceptron(n_iter=40, eta0=0.1, random_state=1)
ppn.fit(X_train_std, y_train)
```

Check the accuracy of test set

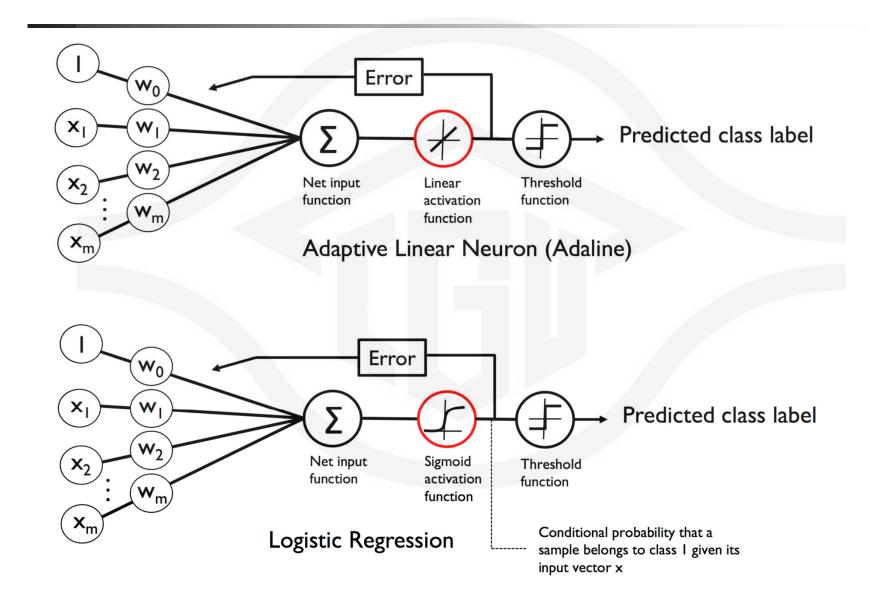
```
y_pred = ppn.predict(X_test_std)
print('Misclassified samples: %d' % (y_test !=
y_pred).sum())
print('Accuracy: %.2f' % accuracy_score(y_test, y_pred))
```

Logistic Regression

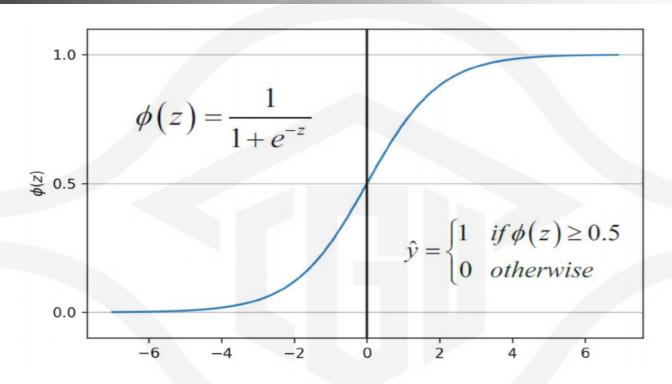
 Logistic Regression is actually a very simple Neural Network



Compare with Adaline



Activation Function with Probability



• Link $\phi(z)$ to Bernoulli's PDF P(y | x, w)

$$P(y|x,w) = p^{y}(1-p)^{1-y} \begin{cases} p, if \ y = 1 \\ 1-p, if \ y = 0 \end{cases}$$

Logistic Regression Characteristic

Replace with the sigmoid function

```
def activation(self, z):
    """Compute logistic sigmoid activation"""
    return 1. / (1. + np.exp(-z))
```

Prediction range constrained between 0 to 1

```
def predict(self, X):
    """Return class label after unit step"""
    return np.where(self.activation(self.net_input(X))
>= 0.5, 1, 0)
```

Maximum Likelihood Estimation

 If training examples were identically independently distributed (IID), then maximum likelihood estimation (MLE) means to find the parameters of the model with given training data

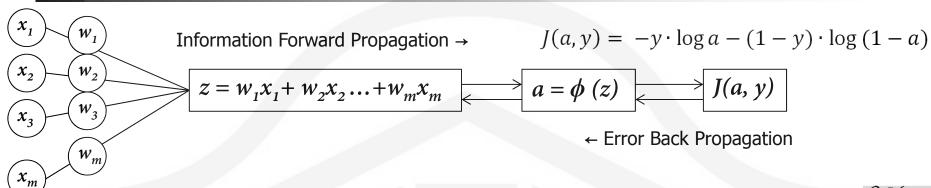
$$L(w) = P(y \mid x; w) = \prod_{i=1}^{n} P(y^{(i)} \mid x^{(i)}; w) = \prod_{i=1}^{n} (\phi(z^{(i)}))^{y^{(i)}} (1 - \phi(z^{(i)}))^{1 - y^{(i)}}$$

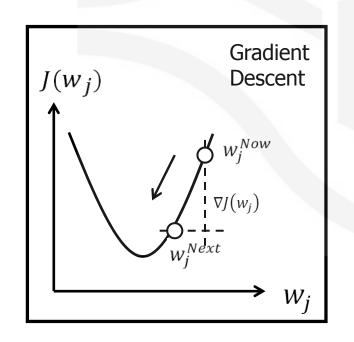
$$l(\mathbf{w}) = \log L(\mathbf{w}) = \sum_{i=1}^{n} \left[y^{(i)} \log \left(\phi\left(z^{(i)}\right) \right) + \left(1 - y^{(i)}\right) \log \left(1 - \phi\left(z^{(i)}\right) \right) \right]$$

Negative log-likelihood as the cost function

$$J(w) = \sum_{i=1}^{n} \left[-y^{(i)} \log \left(\phi(z^{(i)}) \right) - \left(1 - y^{(i)} \right) \log \left(1 - \phi(z^{(i)}) \right) \right]$$

Learning Phase





$$\frac{\partial z}{\partial w_{j}} \times \frac{\partial a}{\partial z} \times \frac{\partial J}{\partial a} = \frac{\partial J(w_{j})}{\partial w_{j}}$$

$$\downarrow \qquad \qquad \downarrow \qquad \qquad$$

$$w_j^{Next} = w_j^{Now} - \eta \cdot \nabla J(w_j)$$

About Cost Function

Logistic regression or conditional log-likelihood cost **function** $(-\log P(y|x))$ worked much better than the quadratic cost(squared errors) which was traditionally used to train feedforward neural networks for classification problems.

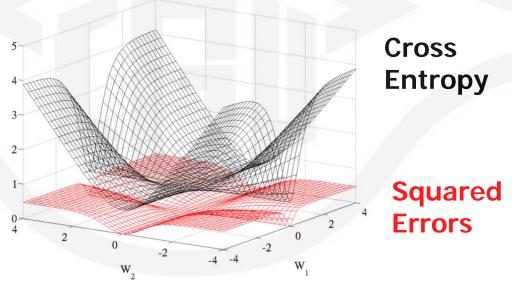


Figure 5: Cross entropy (black, surface on top) and quadratic (red, bottom surface) cost as a function of two

Logistic Regression Training Algorithm

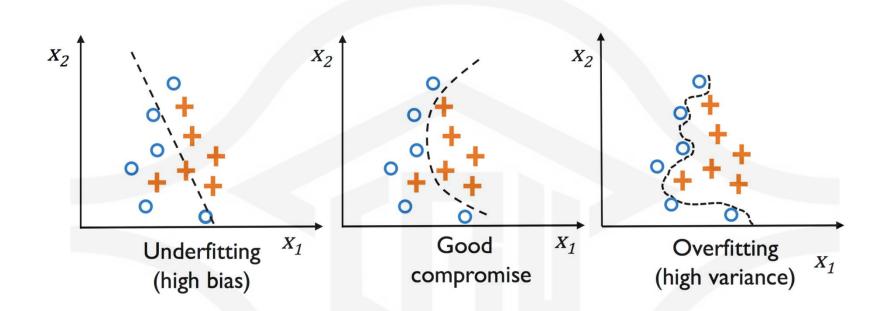
Implement

```
def fit(self, X, y):
  rgen = np.random.RandomState(self.random state)
  self.w_ = rgen.normal(loc=0.0, scale=0.01, size=1 + X.shape[1])
  self.cost = []
  for _ in range(self.n_iter):
      net input = self.net input(X)
      output = self.activation(net input)
      errors = (y - output)
      self.w [1:] += self.eta * X.T.dot(errors)
      self.w [0] += self.eta * errors.sum()
      # note that we compute the logistic `cost` now
      # instead of the sum of squared errors cost
      cost = -y.dot(np.log(output)) - ((1 - y).dot(np.log(1 - output)))
      self.cost .append(cost)
  return self
```

Logistic Regression via sklearn

Initialize the object and training

Tackling Overfitting via Regularization



L2 regularization

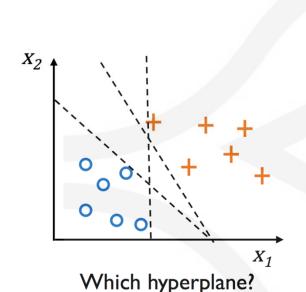
$$J(w) = \sum_{i=1}^{n} \left[-y^{(i)} \log \left(\phi(z^{(i)}) \right) - \left(1 - y^{(i)} \right) \log \left(1 - \phi(z^{(i)}) \right) \right] + \frac{\lambda}{2} ||w||^{2}$$

Regularization Parameter

• C is regularization parameter λ 's inverse

```
weights, params = [], []
for c in np.arange(-5, 5):
    lr = LogisticRegression(C=10.**c, random_state=1)
    lr.fit(X train std, y train)
   weights.append(lr.coef [2])
    params.append(10.**c)
weights = np.array(weights)
plt.plot(params, weights[:, 0],
         label='petal length')
plt.plot(params, weights[:, 1], linestyle='--',
         label='petal width')
plt.ylabel('weight coefficient')
plt.xlabel('C')
plt.legend(loc='upper left')
plt.xscale('log')
plt.show()
```

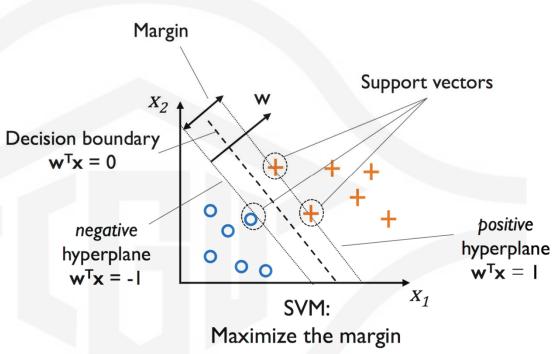
Support Vector Machine



$$w_0 + \boldsymbol{w}^T \boldsymbol{x}_{pos} = 1 \qquad (1)$$

$$w_0 + \boldsymbol{w}^T \boldsymbol{x}_{neg} = -1 \quad (2)$$

$$\Rightarrow \mathbf{w}^T \left(\mathbf{x}_{pos} - \mathbf{x}_{neg} \right) = 2$$



$$\|\mathbf{w}\| = \sqrt{\sum_{j=1}^{m} w_j^2}$$

$$\frac{\boldsymbol{w}^{T}\left(\boldsymbol{x}_{pos} - \boldsymbol{x}_{neg}\right)}{\left\|\boldsymbol{w}\right\|} = \frac{2}{\left\|\boldsymbol{w}\right\|}$$

$$w_0 + \mathbf{w}^T \mathbf{x}^{(i)} \ge 1 \text{ if } y^{(i)} = 1$$

$$w_0 + \mathbf{w}^T \mathbf{x}^{(i)} \le -1 \text{ if } y^{(i)} = -1$$

for
$$i = 1 \dots N$$

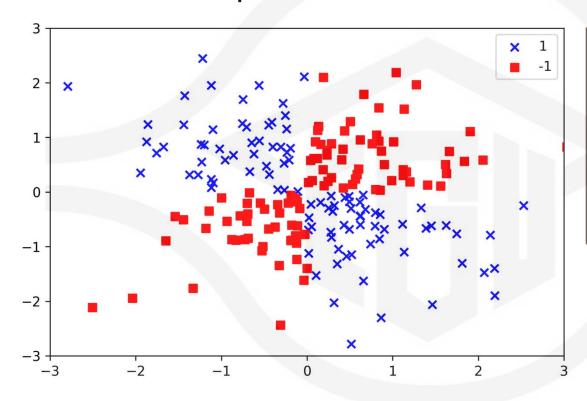
Linear Support Vector Machine via sklearn

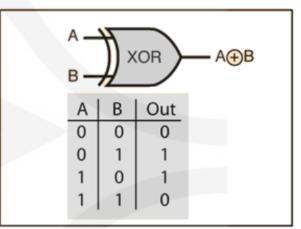
Initialize the object and training

```
from sklearn.svm import SVC
svm = SVC(kernel='linear', C=1.0, random state=1)
svm = SVC(kernel='rbf', random_state=1, gamma=0.10,
C=10.0
svm.fit(X_train_std, y_train)
plot_decision_regions(X_combined_std,
                      y combined,
                      classifier=svm,
                      test_idx=range(105, 150))
plt.xlabel('petal length [standardized]')
plt.ylabel('petal width [standardized]')
plt.legend(loc='upper left')
plt.tight layout()
plt.show()
```

Non-Linear Problems

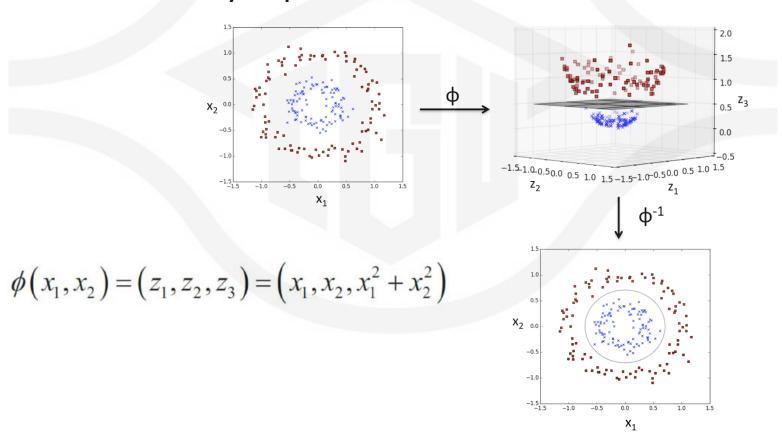
Exclusive-OR problems





Kernel Methods

 Create nonlinear combinations of original features to project them onto a higher-dimensional space where it becomes linearly separable.



Separating Hyperplanes in High-dimensional Space

- Increase features' dimension is computationally expensive especially if dealing with high-dimensional data
- Kernel trick
 - Replace the dot product x(i)T x(j) by K(x(i),x(j))
- Radial Basis Function (RBF) kernel or Gaussian kernel
 - Similarity function between a pair of samples.
 - A range between 1 (for exactly similar samples) and 0 (for very dissimilar samples)

$$\mathcal{K}\left(\boldsymbol{x}^{(i)}, \boldsymbol{x}^{(j)}\right) = \exp\left(-\gamma \left\|\boldsymbol{x}^{(i)} - \boldsymbol{x}^{(j)}\right\|^{2}\right)$$

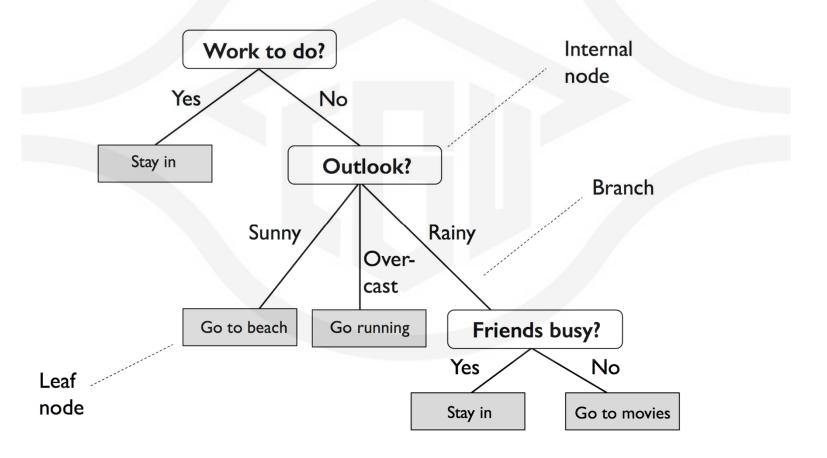
Using the Kernel Trick

RBF kernel

```
np.random.seed(1)
X_xor = np.random.randn(200, 2)
y xor = np.logical_xor(X_xor[:, 0] > 0,
                       X_{xor}[:, 1] > 0)
y_xor = np.where(y_xor, 1, -1)
svm = SVC(kernel='rbf', random_state=1, gamma=0.10,
C=10.0)
svm.fit(X_xor, y_xor)
plot_decision_regions(X_xor, y_xor,
                      classifier=svm)
plt.legend(loc='upper left')
plt.tight_layout()
plt.show()
```

Decision Tree

Interpretability: make a decision based on asking a series of questions



Information Gain

- Using the decision algorithm, we start at the tree root and split the data on the feature that results in the largest Information Gain (IG)
- Maximizing information gain

$$IG\left(D_{p},f\right) = I\left(D_{p}\right) - \frac{N_{left}}{N_{p}}I\left(D_{left}\right) - \frac{N_{right}}{N_{p}}I\left(D_{right}\right)$$

Impurity Measure by Entropy (IH)

$$I_{H}(t) = -\sum_{i=1}^{c} p(i | t) \log_{2} p(i | t)$$

- $\mathbf{p}(\mathbf{i}|\mathbf{t})$: proportion of samples that belong to class i for a node \mathbf{t}
 - The entropy is 0 if all samples belong to the same class
 - The entropy is maximal if a uniform class distribution
 - For example, entropy=0 if p(i=1|t)=1, p(i=0|t|)=0; entropy=1 if p(i=1|t)=0.5, p(i=0|t|)=0.5.
- The entropy criterion attempts to maximize the mutual information in the tree

Gini Impurity (IG)

$$I_G(t) = \sum_{i=1}^{c} p(i \mid t) (1 - p(i \mid t)) = 1 - \sum_{i=1}^{c} p(i \mid t)^2$$

 Gini impurity is maximal if the classes are perfectly mixed, for example, in a binary class setting (c = 2):

$$I_G(t) = 1 - \sum_{i=1}^{c} 0.5^2 = 0.5$$

Impurity Measure by Classification Error (IE)

$$I_E = 1 - \max \{ p(i | t) \}$$

 A useful criterion for pruning but not recommended for growing a decision tree, since it is less sensitive to changes in the class probabilities of the nodes.

Decision Tree via sklearn

```
from sklearn.tree import DecisionTreeClassifier
tree = DecisionTreeClassifier(criterion='gini',
                              max_depth=4,
                              random_state=1)
tree.fit(X_train, y_train)
X_combined = np.vstack((X_train, X_test))
y_combined = np.hstack((y_train, y_test))
plot_decision_regions(X_combined, y_combined,
                      classifier=tree,
test_idx=range(105, 150))
plt.xlabel('petal length [cm]')
plt.ylabel('petal width [cm]')
plt.legend(loc='upper left')
plt.tight_layout()
plt.show()
```

Max Depth

```
petal width <= 0.75
gini = 0.667
samples = 105
value = [35, 35, 35]
class = Setosa
```

gini = 0.0 samples = 35 value = [35, 0, 0] class = Setosa

True

gini = 0.5 samples = 70 value = [0, 35, 35] class = Versicolor

False

Multiple Decision Trees to Random Forests

Random forest algorithm

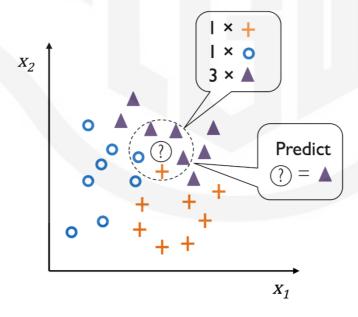
- Draw a random bootstrap sample (randomly choose n samples from the training set).
- Grow a decision tree from the bootstrap sample. At each node:
 - Randomly select d features.
 - Split the node using the feature that provides the best split (maximal information gain).
- Repeat the steps 1-2 k times.
- Aggregate the prediction by each tree to assign the class label by majority vote.

Random Forests via sklearn

```
from sklearn.ensemble import RandomForestClassifier
forest = RandomForestClassifier(criterion='gini',
                                 n_estimators=25,
                                 random_state=1,
                                 n jobs=2
forest.fit(X_train, y_train)
plot_decision_regions(X_combined, y_combined,
                      classifier=forest,
test_idx=range(105, 150))
plt.xlabel('petal length [cm]')
plt.ylabel('petal width [cm]')
plt.legend(loc='upper left')
plt.tight_layout()
plt.show()
```

K-Nearest Neighbors

- It doesn't learn a discriminative function from the training data, but memorizes the training dataset instead
 - Choose the number of k and a distance metric.
 - Find the k-nearest neighbors of the sample to classify.
 - Assign the class label by majority vote.



Minkowski distance

 Minkowski distance is typically used with p being 1 or 2, which correspond to the Manhattan distance and the Euclidean distance, respectively.

$$D\left(X,Y
ight) = \left(\sum_{i=1}^{n}\left|x_{i}-y_{i}
ight|^{p}
ight)^{1/p}$$

K-Nearest Neighbors via sklearn

```
from sklearn.neighbors import KNeighborsClassifier
knn = KNeighborsClassifier(n neighbors=5,
                           p=2,
                           metric='minkowski')
knn.fit(X_train_std, y_train)
plot_decision_regions(X_combined_std, y_combined,
                      classifier=knn,
test idx=range(105, 150))
plt.xlabel('petal length [standardized]')
plt.ylabel('petal width [standardized]')
plt.legend(loc='upper left')
plt.tight_layout()
plt.show()
```

Reference

 Sebastian Raschka, Vahid Mirjalili. Python Machine Learning: Machine Learning and Deep Learning with Python, scikit-learn, and TensorFlow. Second Edition. Packt Publishing, 2017.