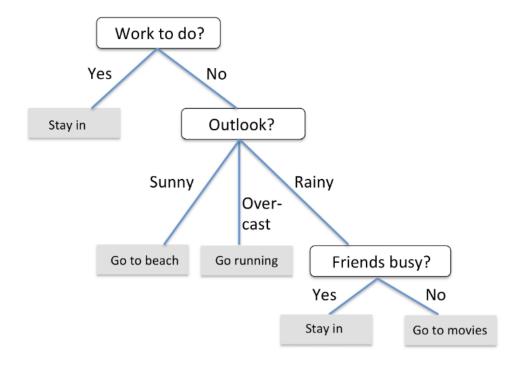


Decision Trees, Bayesian Classifier

Machine Learning

- What is the Decision Tree Learning?
 - In computer science, Decision tree learning uses a decision tree(as a predictive model) to go from observations about an item(represented in the branches) to conclusions about the item's target value(represented in the leaves)





- Maximizing information gain
 - Objective function

$$IG(D_p, f) = I(D_p) - \sum_{j=1}^{m} \frac{N_j}{N_p} I(D_j)$$

f: features for classification

 D_p : Parent node

 D_i : jth child node

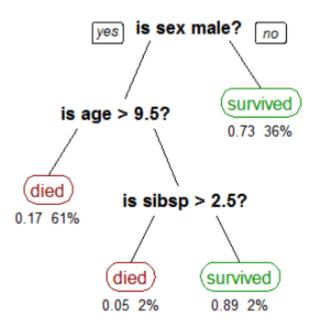
I: Impurity

 N_p : Number of samples in parent node

 N_j : Number of samples in jth child node

- Maximizing information gain
 - Objective function for binary decision tree

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



Most libraries (Including Scikit-learn) use Binary Decision Tree to reduce Search Space

- Maximizing information gain
 - Three Impurity methods
 - Entropy(I_H)

$$I_H(t) = -\sum_{i=1}^{c} p(i|t)log_2 p(i|t)$$

• Gini impurity(I_G)

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

- Maximizing information gain
 - Three Impurity methods
 - Entropy(I_H)

$$I_H(t) = -\sum_{i=1}^{c} p(i|t)log_2 p(i|t)$$

p(i|t): the proportion of the samples that belong to class c for particular node t

Ex) binary classification(c = 2)

$$p(i = 0|t) = 0(or 1)$$

 $\rightarrow I_H(t) = 0 \text{ (min)}$

$$p(i = 0|t) = \frac{1}{2}$$

$$\rightarrow I_H(t) = 1 \text{ (max)}$$

- Maximizing information gain
 - Three Impurity methods
 - Gini impurity(I_G)

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

p(i|t): the proportion of the samples that belong to class c for particular node t

Ex) binary classification(c = 2)

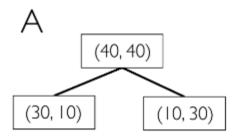
$$p(i = 0|t) = 0$$

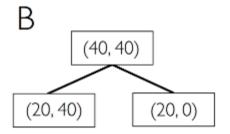
 $\rightarrow I_G(t) = 1 - 1^2 - 0^2 = 0$ (min)

$$p(i = 0|t) = \frac{1}{2}$$

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

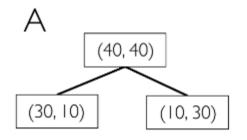
- Maximizing information gain
 - Three Impurity methods

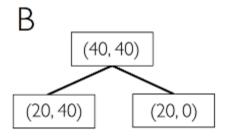




- Entropy(I_H)
- Gini impurity(I_G)

- Maximizing information gain
 - Three Impurity methods





• Entropy(I_H)

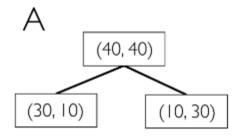
$$I_{H}(D_{P}) = -(0.5 \cdot log_{2}(0.5) + 0.5 \cdot log_{2}(0.5)) = 1$$

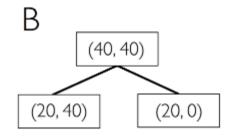
$$A: I_{H}(D_{left}) = -(0.75 \cdot log_{2}(0.75) + 0.25 \cdot log_{2}(0.25)) = 0.81$$

$$A: I_{H}(D_{right}) = -(0.25 \cdot log_{2}(0.25) + 0.75 \cdot log_{2}(0.75)) = 0.81$$

$$A: IG_{H} = 1 - \frac{4}{8} \cdot 0.81 - \frac{4}{8} \cdot 0.81 = 0.19$$

- Maximizing information gain
 - Three Impurity methods





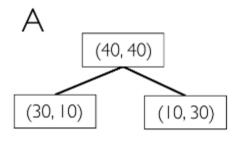
• Entropy(I_H)

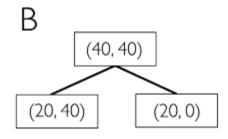
$$B: I_{H}(D_{left}) = -(\frac{2}{6} \cdot log_{2}\left(\frac{2}{6}\right) + \frac{4}{6} \cdot log_{2}\left(\frac{4}{6}\right)) = 0.92$$

$$B: I_{H}(D_{right}) = 0$$

$$B: IG_{H} = 1 - \frac{6}{8} \cdot 0.92 - \frac{2}{8} \cdot 0 = 0.31$$
Using Entropy(I_{H}), Scenario B will be selected!(0.19 < 0.31)

- Maximizing information gain
 - Three Impurity methods



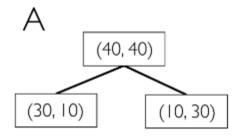


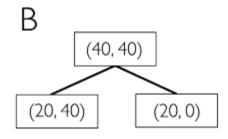
• Gini impurity(I_G)

$$I_G(D_P) = 1 - (0.5^2 + 0.5^2) = 0.5$$

 $A: I_G(D_{left}) = 1 - (0.75^2 + 0.25^2) = 0.375$
 $A: I_G(D_{right}) = 1 - (0.25^2 + 0.75^2) = 0.375$
 $A: IG_G = 0.5 - \frac{4}{8}0.375 - \frac{4}{8}0.375 = 0.125$

- Maximizing information gain
 - Three Impurity methods





• Gini impurity(I_G)

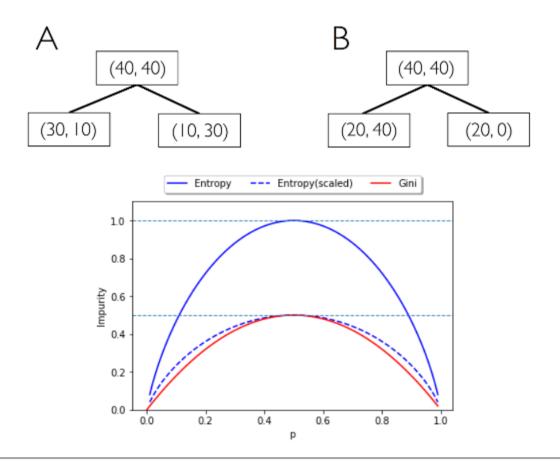
$$B: I_G(D_{left}) = 1 - ((2/6)^2 + (4/6)^2) = \frac{4}{9}$$

$$B: I_G(D_{right}) = 1 - (1^2 + 0^2) = 0$$

$$B: IG_G = 0.5 - \frac{6}{8} \cdot \frac{4}{9} - \frac{2}{8} \cdot 0 = \frac{1}{6}$$

Using Gini impurity(I_G), Scenario B will be selected!(0.125 < $\frac{1}{6}$)

- Maximizing information gain
 - Three Impurity methods



Load Iris Dataset

```
from sklearn import datasets
import numpy as np
iris = datasets.load_iris()
# use features 2 and 3 only
X = iris.data[:, [2, 3]]
y = iris.target
print('Class labels:', np.unique(y))
Class labels: [0 1 2]
```

Splitting data into 70% training data & 30% test data

```
from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(
X, y, test_size=0.3, random_state=1, stratify=y)

print('Labels counts in y: [50 50 50]
Labels counts in y_train: [35 35 35]
Labels counts in y_test: [15 15 15]

print('Labels counts in y_train:', np.bincount(y))
print('Labels counts in y_test:', np.bincount(y_test))
```

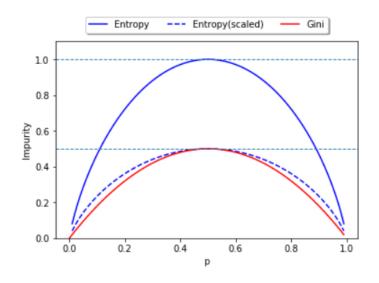
Entropy and Gini Impurity

```
def gini(p):
    return p * (1 - p) + (1 - p) * (1 - (1 - p))
def entropy(p):
    entropy(p): return - p * np.log2(p) - (1 - p) * np.log2((1 - p)) \longrightarrow I_H(t) = -\sum_{i=1}^c p(i|t)log_2p(i|t)
```

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

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

$$I_H(t) = -\sum_{i=1}^{c} p(i|t)log_2 p(i|t)$$



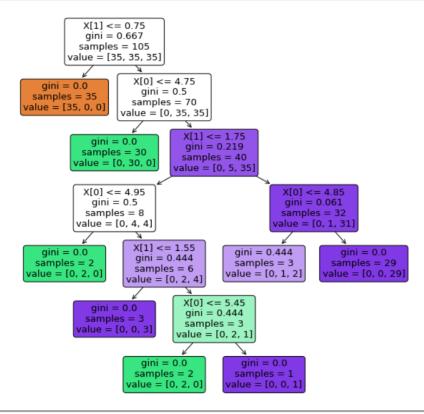
Learning Decision Tree

Model evaluation

```
X_test[10:15]
array([[1.5, 0.4],
      [4.9, 1.8],
      [1.4, 0.2],
      [3.3, 1.],
      [1.4, 0.2]]
y test[10:15]
array([0, 2, 0, 1, 0])
# predict class of X_test[10] ~ X_test[14]
tree.predict(X test[10:15])
array([0, 2, 0, 1, 0])
# Compute train accuracy
acc = tree.score(X train, y train)
print("Train Accuracy : ", acc)
Train Accuracy: 0.9904761904761905
# Compute test accuracy
acc = tree.score(X test, y test)
print("Test Accuracy : ", acc)
```

Visualizing the model

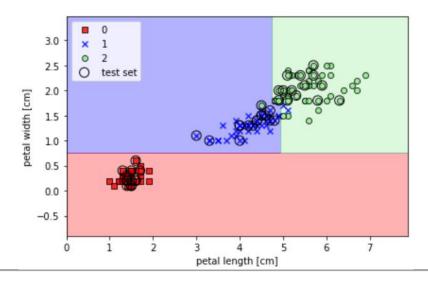
```
from sklearn.tree import plot_tree #scikit-learn >= 22.0
# plot the tree
plot_tree(tree, filled=True, rounded=True)
```



Plotting the decision boundary

```
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()
```



Building Decision Tree with max depth

Model evaluation

```
# Compute train accuracy
acc = tree.score(X_train, y_train)
print("Train Accuracy : ", acc)
Train Accuracy : 0.9523809523809523
# Compute test accuracy
acc = tree.score(X_test, y_test)
print("Test Accuracy : ", acc)
Test Accuracy : 0.9555555555555556
```

Visualizing the model

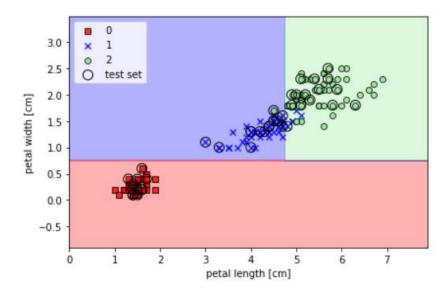
```
from sklearn.tree import plot_tree #scikit-learn >= 22.0
# plot the tree
plot_tree(tree, filled=True, rounded=True)
```

```
X[1] \le 0.75
          entropy = 0.667
          samples = 105
        value = [35, 35, 35]
                     X[0] <= 4.75
 entropy = 0.0
                     entropy = 0.5
 samples = 35
                     samples = 70
/alue = [35, 0, 0]
                   value = [0, 35, 35]
           entropv = 0.0
                              entropy = 0.219
           samples = 30
                               samples = 40
         value = [0, 30, 0]
                              value = [0, 5, 35]
```

Plotting the decision boundary

```
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()
```



Naïve Bayesian Classifiers

- What is the Naïve Bayesian Classifiers?
 - In machine learning, Naïve Bayesian Classifiers are a family of simple "probabilistic classifiers" based on applying Bayes' theorem with strong (Naïve) independence assumptions between the features
 - Probabilistic model
 - Naïve Bayes is a conditional probability model : given a problem instance to be classified, represented by vector $x = (x_1, x_2, ..., x_n)$ representing some n features(independent), it assigns to this instance probabilities $p(C_k|x_1,...,x_n)$

for each of K possible outcomes of classes C_k

Bayesian probability terminology

$$posterior = \frac{prior \times likelihood}{evidence} \qquad p(C_k|x) = \frac{p(C_k)p(x|C_k)}{p(x)}$$

Naïve Bayesian Classifiers

- Constructing a classifier from the probability model
 - Bayesian Reasoning

$$p(x_i|x_{i+1},...,x_n,C_k) = p(x_i|C_k)$$

$$p(C_k|x_1,...,x_n) = \frac{1}{Z}p(C_k)\prod_{i=1}^n p(x_i|C_k)$$

$$where Z = p(x) = \sum_k p(C_k)p(x|C_k)$$

Thus, the joint model can be expressed as

$$p(C_k|x_1, ..., x_n) \propto p(C_k, x_1, ..., x_n)$$

$$= p(C_k)p(x_1|C_k)p(x_2|C_k)p(x_3|C_k) ... = p(C_k) \prod_{i=1}^n p(x_i|C_k)$$

Prediction

$$\hat{y} = \operatorname*{argmax}_{k \in \{1, \dots, K\}} p(C_k) \prod_{i=1}^{n} p(x_i | C_k)$$



Load Iris Dataset

```
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import numpy as np
iris = datasets.load_iris()
# use features 2 and 3 only
X = iris.data[:, [2, 3]]
y = iris.target
print('Class labels:', np.unique(y))
Class labels: [0 1 2]
```

Splitting data into 70% training data & 30% test data

```
from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(
X, y, test_size=0.3, random_state=1, stratify=y)

print('Labels counts in y: [50 50 50]
Labels counts in y_train: [35 35 35]
Labels counts in y_test: [15 15 15]

print('Labels counts in y_train:', np.bincount(y))
print('Labels counts in y_test:', np.bincount(y_test))
```

Learning Gaussian Naïve Bayes Classifier

```
from sklearn.naive_bayes import GaussianNB

gnb = GaussianNB()
gnb.fit(X_train, y_train)

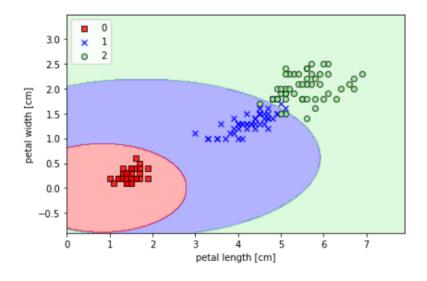
GaussianNB(priors=None, var_smoothing=1e-09)
```

Model evaluation - (1)

Model evaluation –(2)

Plotting the decision boundary

```
from sklearn.naive_bayes import GaussianNB
X_combined = np.vstack((X_train, X_test))
y_combined = np.hstack((y_train, y_test))
plot_decision_regions(X_combined, y_combined,
classifier=gnb)
plt.xlabel('petal length [cm]')
plt.ylabel('petal width [cm]')
plt.legend(loc='upper left')
plt.tight_layout()
plt.show()
```



Submit

- To make sure if you have completed this practice,
 Submit your practice file(Week06_givencode.ipynb) to e-class.
- Deadline : tomorrow 11:59pm
- Modify your ipynb file name as "Week06_StudentNum_Name.ipynb"
 Ex) Week06_2020123456_홍길동.ipynb
- You can upload this file without taking the quiz, but homework will be provided like a quiz every three weeks, so it is recommended to take the quiz as well.

Quiz 1 : Decision Tree

- Learn decision tree model for Heart Diseases classification using heart_disease.csv dataset
 - Dataset information : https://archive.ics.uci.edu/ml/datasets/heart+Disease
 - 297 samples, all 13 features
 - Label: The feature 'num' refers to the presence of heart disease in the patient(from 0 to 4). We will convert values 2, 3, 4 to 1(to do binary classification)
 - Find the smallest tree among the trees that show highest test accuracy
 - Visualize the tree, and compute the accuracies
 - Predict the class of following data [[52, 2, 5, 135, 250, 0, 3, 180, 1, 0, 1, 0, 2]]

Quiz 2 : Naïve Bayesian Classifier

- Predicting presence of Heart Diseases using Bayesian
 Classifier
 - Dataset information : https://archive.ics.uci.edu/ml/datasets/heart+Disease
 - Use the same training and test dataset of Quiz 1
 - Compute the accuracies of the classifier
 - Predict the class of following data with probability [[52, 2, 5, 135, 250, 0, 3, 180, 1, 0, 1, 0, 2]]