```
In [1]: import numpy as np
    import pandas as pd
    from sklearn import tree
    from sklearn.model_selection import GridSearchCV
    from matplotlib import pyplot as plt
    import graphviz
```

```
In [2]: df = pd.read_csv('Data/Iris/Iris-cleaned.csv')
    df.head(3)
```

#### Out[2]:

	sepal length	sepal width	petal length	petal width	species
0	5.1	3.5	1.4	0.2	setosa
1	4.9	3.0	1.4	0.2	setosa
2	4.7	3.2	1.3	0.2	setosa

### (a) Preprocessing:

Randomize the order of the records using the command below.

```
In [5]: df = df.sample(frac=1.0,random_state=0)
```

## (b) Fit a depth 2 decision tree for predicting iris species. Define classes as shown below

```
In [6]: class_names = df.species.cat.categories
class_names

Out[6]: Index(['setosa', 'versicolor', 'virginica'], dtype='object')

In [7]: attributes = df.drop(['species'], axis = 1) #continuous -> one hot encode
    attributes = pd.get_dummies(attributes)
    target = df.species
```

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```
In [8]: Tree = tree.DecisionTreeClassifier()
 In [9]: Tree = tree.DecisionTreeClassifier(max_depth = 2)
         Tree.fit(attributes, target)
Out[9]: DecisionTreeClassifier(max_depth=2)
In [10]: dot data = tree.export graphviz(Tree,
                                        out file=None,
                                        feature_names=attributes.columns,
                                        class names = df.species.cat.categories,
                                        proportion=True,
                                        impurity=False,
                                        filled=True,
                                        rounded=True)
         graph = graphviz.Source(dot_data)
         graph.render('Iris')
         graph
Out[10]:
                           petal width <= 0.8
                           samples = 100.0%
                     value = [0.333, 0.333, 0.333]
                             class = setosa
                                            False
                       True
                                         petal width <= 1.75
             samples = 33.3%
                                          samples = 66.7%
           value = [1.0, 0.0, 0.0]
                                        value = [0.0, 0.5, 0.5]
               class = setosa
                                          class = versicolor
                         samples = 36.0%
                                                           samples = 30.7\%
                    value = [0.0, 0.907, 0.093]
                                                      value = [0.0, 0.022, 0.978]
                                                            class = virginica
                         class = versicolor
```

(c) Compute species group averages for the attributes and use these attribute averages to check that your decision tree is reasonable.

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6.588

```
In [11]: df_mean = df.groupby('species').mean()
    df_mean
```

5.552

2.026

sepal length sepal width petal length petal width

#### Out[11]:

species				
setosa	5.006	3.418	1.464	0.244
versicolor	5.936	2.770	4.260	1.326

2.974

The average petal length of setosa is significantly smaller than others. Most setosa have petal length <= 2.45. Tt is reasonable to use petal width as an attribute to categorize because most versicolors have petal width <= 1.75 and most virginicas have petal width > 1.75.

```
In [12]: Tree.predict(df_mean)==df_mean.index
Out[12]: array([ True, True, True])
```

So the te deicison tree is reasonable

virginica

# (d) Use cross-validation and the hyper-parameters shown below to determine a tree that minimizes generalization error.

```
In [13]: Tree = tree.DecisionTreeClassifier()
# 'min_samples_split': [5,10,15,20,25,30,35,40,45,50,55,60,65,70] -> output is
40 -> check number around 40
# then: 'min_samples_split': [35,36,37,38,39,40,41,42,43,44,45]
parameters = {'max_depth':[1,2,3,4,5,6,7,8,9,10], 'min_samples_split': [2,3,4,5,6,7,8,9,10]}
TreeCV = GridSearchCV(Tree, param_grid = parameters, n_jobs=-1)
TreeCV.fit(attributes,target)
print(TreeCV.best_params_)
print(TreeCV.best_score_)
print(TreeCV.best_estimator_)

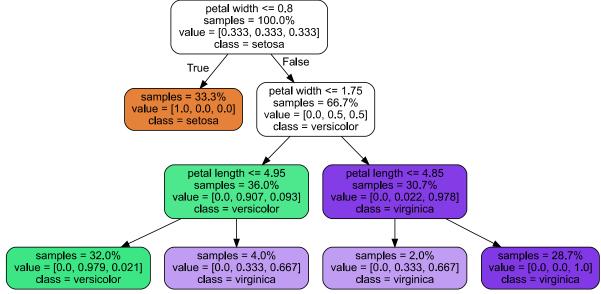
{'max_depth': 3, 'min_samples_split': 2}
0.96
DecisionTreeClassifier(max_depth=3)
```

max depth — maximum depth of the tree.

min samples split — mimimum size of a node that is allowed to be split. (Regulates the complexity of the tree.)

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#### Out[14]:



```
In [15]: #results = pd.DataFrame()
    #results['depth'] = parameters['max_depth']
    #results['valid error'] = 1 - TreeCV.cv_results_['mean_test_score'].mean()
    #results.head()
    #results.plot.line(x='depth',y='valid error')
```