Kayvan Shah Assignment 2 solution

January 31, 2023

1 Assignment 2: Exploratory Data Analysis and K Nearest Neighbors Classification

For this assignment you will perform exploratory data analysis to visualize Fisher's Iris dataset using Scikit Learn. And, you will explore the bias/variance trade-off by applying k-nearest neighbors classification to the Iris dataset and varying the hyperparameter k.

Documentation for Scikit Learn: + The top level documenation page is here: https://scikit-learn.org/stable/index.html + The API for the KNearestNeighborsClassifier is here: https://scikit-learn.org/stable/modules/generated/sklearn.neighbors.KNeighborsClassifier.html#sklearn.neighbors.KNeighborsClassifier is here: https://scikit-learn.org/stable/modules/neighbors.html#classification + Scikit Learn provides many Jupyter notebook examples on how use the toolkit. These Jupyter notebook examples can be run on MyBinder: https://scikit-learn.org/stable/auto_examples/index.html

For more information about the Iris dataset, see this page https://en.wikipedia.org/wiki/Iris_flower_data_set.

```
[]: | %%bash | pip install -U scikit-learn |
```

Looking in indexes: https://pypi.org/simple, https://us-python.pkg.dev/colab-wheels/public/simple/
Requirement already satisfied: scikit-learn in /usr/local/lib/python3.8/dist-packages (1.2.1)
Requirement already satisfied: scipy>=1.3.2 in /usr/local/lib/python3.8/dist-packages (from scikit-learn) (1.7.3)
Requirement already satisfied: numpy>=1.17.3 in /usr/local/lib/python3.8/dist-packages (from scikit-learn) (1.21.6)
Requirement already satisfied: threadpoolctl>=2.0.0 in
/usr/local/lib/python3.8/dist-packages (from scikit-learn) (3.1.0)
Requirement already satisfied: joblib>=1.1.1 in /usr/local/lib/python3.8/dist-packages (from scikit-learn) (1.2.0)

```
[]: import matplotlib.pyplot as plt
from mpl_toolkits.mplot3d import Axes3D
from sklearn import datasets
from sklearn import neighbors
from sklearn.model_selection import train_test_split
```

```
from pandas import DataFrame
    Load Iris dataset
[]: iris = datasets.load iris()
     X = iris.data
     y = iris.target
[]: print("Number of instances in the iris dataset:", X.shape[0])
     print("Number of features in the iris dataset:", X.shape[1])
     print("The dimension of the data matrix X is", X.shape)
    Number of instances in the iris dataset: 150
    Number of features in the iris dataset: 4
    The dimension of the data matrix X is (150, 4)
[]: X
[]: array([[5.1, 3.5, 1.4, 0.2],
            [4.9, 3., 1.4, 0.2],
            [4.7, 3.2, 1.3, 0.2],
            [4.6, 3.1, 1.5, 0.2],
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            [4.8, 3., 1.4, 0.1],
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            [5.8, 4., 1.2, 0.2],
            [5.7, 4.4, 1.5, 0.4],
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```

```
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[6.4, 2.9, 4.3, 1.3],
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[6.8, 2.8, 4.8, 1.4],
```

```
[6.7, 3., 5., 1.7],
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[5.5, 2.4, 3.7, 1.],
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[6.8, 3., 5.5, 2.1],
[5.7, 2.5, 5., 2.],
[5.8, 2.8, 5.1, 2.4],
[6.4, 3.2, 5.3, 2.3],
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[5.6, 2.8, 4.9, 2.],
[7.7, 2.8, 6.7, 2.],
[6.3, 2.7, 4.9, 1.8],
```

```
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[6.2, 2.8, 4.8, 1.8],
[6.1, 3., 4.9, 1.8],
[6.4, 2.8, 5.6, 2.1],
[7.2, 3., 5.8, 1.6],
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[6.3, 2.8, 5.1, 1.5],
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[7.7, 3., 6.1, 2.3],
[6.3, 3.4, 5.6, 2.4],
[6.4, 3.1, 5.5, 1.8],
[6., 3., 4.8, 1.8],
[6.9, 3.1, 5.4, 2.1],
[6.7, 3.1, 5.6, 2.4],
[6.9, 3.1, 5.1, 2.3],
[5.8, 2.7, 5.1, 1.9],
[6.8, 3.2, 5.9, 2.3],
[6.7, 3.3, 5.7, 2.5],
[6.7, 3., 5.2, 2.3],
[6.3, 2.5, 5., 1.9],
[6.5, 3., 5.2, 2.],
[6.2, 3.4, 5.4, 2.3],
[5.9, 3., 5.1, 1.8]
```

The y vector length is 150. It has three unique values: 0, 1 and 2. Each value represents a species of iris flower.

```
[]: ['DESCR',
      'data',
      'data_module',
      'feature_names',
      'filename',
      'frame',
      'target',
      'target_names']
[]: iris.target_names
[]: array(['setosa', 'versicolor', 'virginica'], dtype='<U10')
[]: iris.feature_names
[]: ['sepal length (cm)',
      'sepal width (cm)',
      'petal length (cm)',
      'petal width (cm)']
    1.0.1 Extension: Show the summary table of iris data including min, max, median,
           quantiles
[]: # Insert your answer here
     import pandas as pd
     iris_df = pd.DataFrame(iris.data, columns=iris.feature_names)
[]: iris_df.describe()
[]:
            sepal length (cm)
                                sepal width (cm)
                                                  petal length (cm)
                   150.000000
                                      150.000000
                                                          150.000000
     count
    mean
                     5.843333
                                        3.057333
                                                            3.758000
                     0.828066
                                        0.435866
                                                            1.765298
     std
                     4.300000
    min
                                        2.000000
                                                            1.000000
     25%
                     5.100000
                                        2.800000
                                                            1.600000
     50%
                     5.800000
                                        3.000000
                                                            4.350000
     75%
                     6.400000
                                        3.300000
                                                            5.100000
                     7.900000
                                        4.400000
                                                            6.900000
     max
            petal width (cm)
                  150.000000
     count
    mean
                    1.199333
     std
                    0.762238
    min
                    0.100000
     25%
                    0.300000
     50%
                    1.300000
```

```
75% 1.800000 max 2.500000
```

1.1 Part 1Exploratory Data Analysis

1.1.1 Part 1a

Generate scatter plots using each pair of the attributes as axis. You should generate $6 = \binom{4}{2}$ scatter plots.

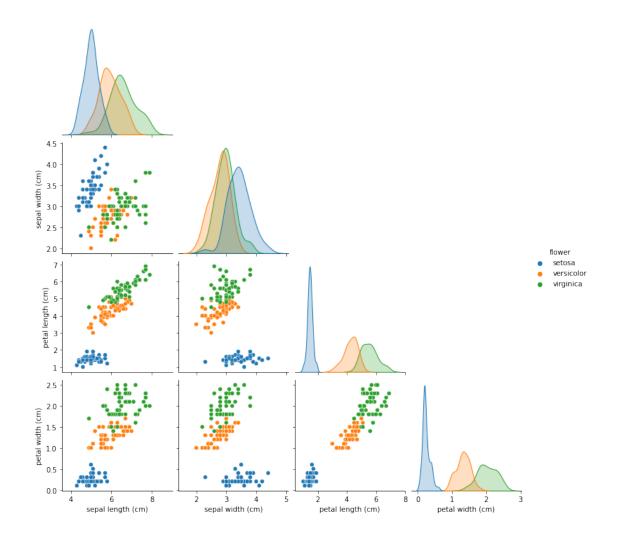
```
sepal length (cm) sepal width (cm) petal length (cm) petal width (cm)
[]:
                      5.1
                                        3.5
                                                            1.4
                                                                              0.2
     1
                      4.9
                                        3.0
                                                            1.4
                                                                              0.2
     2
                      4.7
                                        3.2
                                                           1.3
                                                                              0.2
     3
                      4.6
                                                                              0.2
                                        3.1
                                                            1.5
                      5.0
                                        3.6
                                                            1.4
                                                                              0.2
```

flower

- 0 setosa
- 1 setosa
- 2 setosa
- 3 setosa
- 4 setosa

```
[]: import matplotlib.pyplot as plt
import seaborn as sns
sns.pairplot(iris_df_pp, hue="flower", corner=True)
```

[]: <seaborn.axisgrid.PairGrid at 0x7f45a73a3cd0>



1.1.2 Extension: Draw a boxplot of sepal length (cm), sepal width (cm), petal length (cm), petal width (cm). Use color to show the different target class.

Some links to help you:

https://seaborn.pydata.org/generated/seaborn.boxplot.html

https://matplotlib.org/stable/api/_as_gen/matplotlib.pyplot.boxplot.html

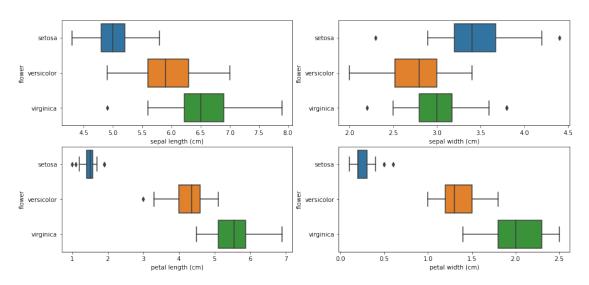
```
[]: # Insert your code ...
# Creating subplots
fig, ax = plt.subplots(2, 2, figsize=(15, 7))
fig.suptitle("Box Plots Iris Features", fontsize=16)

# Box Plot for Sepal Length
sns.boxplot(data=iris_df_pp, x="sepal length (cm)", y="flower", ax=ax[0,0])
# Box Plot for Sepal Length
sns.boxplot(data=iris_df_pp, x="sepal width (cm)", y="flower", ax=ax[0,1])
```

```
# Box Plot for Sepal Length
sns.boxplot(data=iris_df_pp, x="petal length (cm)", y="flower", ax=ax[1,0])
# Box Plot for Sepal Length
sns.boxplot(data=iris_df_pp, x="petal width (cm)", y="flower", ax=ax[1,1])
```

[]: <matplotlib.axes._subplots.AxesSubplot at 0x7f45a6d9c790>

Box Plots Iris Features



1.1.3 Part 1b

If you were to draw linear decision boundaries to separate the classes, which scatter plot do you think will have the least error and which the most?

1.1.4 Insert your 1b answer here

Scatter Plot with:

- Most Error: sepal width (cm) vs sepal length (cm)
- Least Error: sepal width (cm) vs petal length (cm)

1.1.5 Part 1c

Scatter plots using two attributes of the data are equivalent to project the four dimensional data down to two dimensions using axis-parallel projection. Principal component analysis (PCA) is a technique to linearly project the data to lower dimensions that are not necessarily axis-parallel. Use PCA to project the data down to two dimensions.

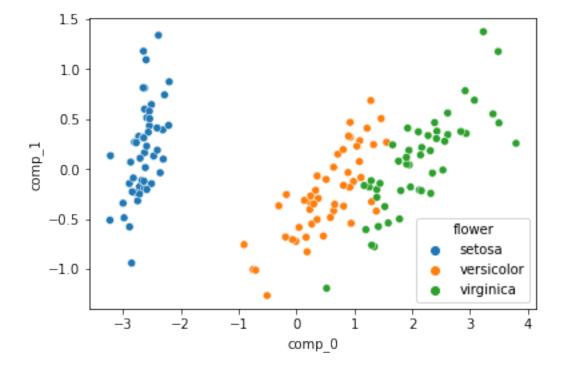
Documentation for PCA: + API https://scikit-learn.org/stable/modules/generated/sklearn.decomposition.PCA.ht + User guide https://scikit-learn.org/stable/modules/decomposition.html#pca

```
[]: comp_0 comp_1 flower 0 -2.684126 0.319397 setosa 1 -2.714142 -0.177001 setosa 2 -2.888991 -0.144949 setosa 3 -2.745343 -0.318299 setosa 4 -2.728717 0.326755 setosa
```

1.1.6 In the case of the Iris dataset, does PCA do a better job of separating the classes?

```
[]: sns.scatterplot(data=data, x="comp_0", y="comp_1", hue="flower")
```

[]: <matplotlib.axes._subplots.AxesSubplot at 0x7f45a6c5a670>



1.1.7 Insert your answer

PCA with n_components=2 does a better job separating the 3 classes with minimal region of overlap between *versicolor* & *virginica*

1.2 Part 2 K Nearest Neighbor

Split the dataset into train set and test set. Use 67 percent of the dataset for training, and use 33 percent for testing.

```
[]: X_train, X_test, y_train, y_test = train_test_split(
     X, y, test_size=0.33, random_state=42
)
```

```
[]: print("Number of instances in the train set:", X_train.shape[0]) print("Number of instances in the test set:", X_test.shape[0])
```

```
Number of instances in the train set: 100 Number of instances in the test set: 50
```

1.2.1 Part 2a Create a KNeibhorsClassifier with n_neighbors = 5. And, train the classifier using the train set.

```
[]: ### Insert you answer here
from sklearn.neighbors import KNeighborsClassifier

model = KNeighborsClassifier(n_neighbors=5)
model.fit(X, y)
```

[]: KNeighborsClassifier()

```
[]: print("Using", model.n_neighbors, "neighbors:")
print("The train accuracy score is:", model.score(X_train, y_train))
print("The test accuracy score is:", model.score(X_test, y_test))
```

Using 5 neighbors:

The train accuracy score is: 0.95 The test accuracy score is: 1.0

1.2.2 Part 2b Tuning hyperparameter k

As we have seen in class, hyperparameter k of the K Nearest Neighbors classification affects the inductive bias. For this part train multiple near neighbor classifier models, store the results in a DataFrame. The plot plot training error and testing error versus N/k, where N=100.

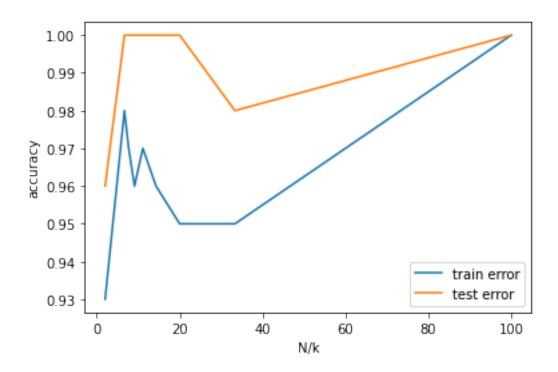
1.2.3 Extension: Use different metric for knn classification.

- 1). Euclidean distance
- 2). Manhattan distance
- 3). Chebyshev distance.

Distance Metrics Documentation: https://scikit-learn.org/stable/modules/generated/sklearn.metrics.pairwise.dist

```
[]: k_list = [1, 3, 5, 7, 9, 11, 13, 15, 50]
     def get_train_test_accuracy(k_list: list=k_list, metric=None):
         train = []
         test = []
         n_k = []
         for k in k_list:
             if metric:
                 model = KNeighborsClassifier(n_neighbors=k, metric=metric)
             else:
                 model = KNeighborsClassifier(n_neighbors=k)
             model.fit(X, y)
             train.append(model.score(X_train, y_train))
             test.append(model.score(X_test, y_test))
             n_k.append(100/k)
         result = pd.DataFrame(
             data = {
                 "N/k": n_k,
                 "train error": train,
                 "test error":test
             },
         return result
[]: ### Insert your code
     # Use the `result` to store the DataFrame
     # euclidean
     result = get_train_test_accuracy(metric="euclidean")
[]: result.plot(x='N/k', y=['train error', 'test error'], ylabel='accuracy')
```

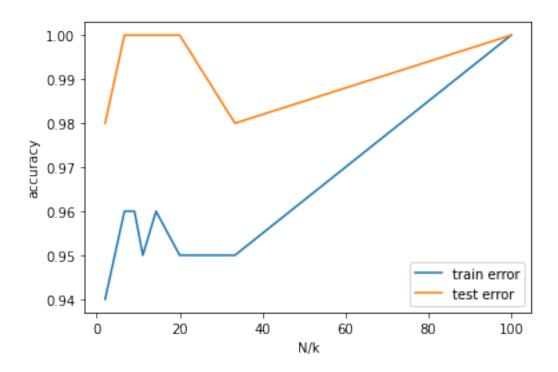
[]: <matplotlib.axes._subplots.AxesSubplot at 0x7f45a6c829a0>



```
[]: ### Insert your code
# Use the `result` to store the DataFrame
# manhattan
result = get_train_test_accuracy(metric="manhattan")

[]: result.plot(x='N/k', y=['train error', 'test error'], ylabel='accuracy')
```

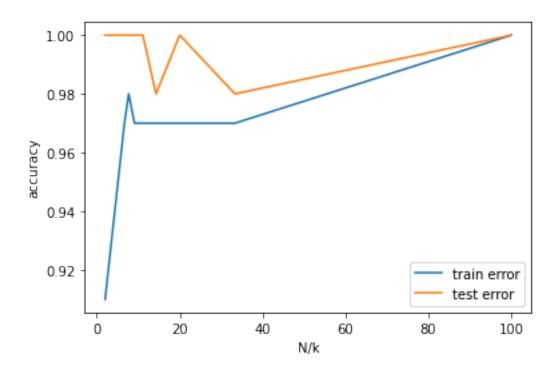
[]: <matplotlib.axes._subplots.AxesSubplot at 0x7f45a6bbef70>



```
[]: ### Insert your code
# Use the `result` to store the DataFrame
# chebyshev
result = get_train_test_accuracy(metric="chebyshev")

[]: result.plot(x='N/k', y=['train error', 'test error'], ylabel='accuracy')
```

[]: <matplotlib.axes._subplots.AxesSubplot at 0x7f45a6b3ed90>



1.2.4 Part 2c Plot decision boundaries of K Nearest Neighbors

Use Scikit Learn's DecisionBoundaryDisplay class to visualize the nearest neighbor boundaries as k is varied.

https://scikit-learn.org/stable/modules/generated/sklearn.inspection.DecisionBoundaryDisplay.html#sklearn.ins

```
[]: k_list = [1, 3, 5, 7, 9, 11, 13, 15, 50]
```

Simplify the problem by using only the first 2 attributes of the dataset

```
def get_pred():
    y = []
    for k in k_list:
        model = KNeighborsClassifier(n_neighbors=k)
        model.fit(X2, iris.target)
        y_pred = np.reshape(model.predict(grid), feat_1.shape)
        y.append(y_pred)
    return y

y = get_pred()
```

```
for i in range(len(k_list)):
    display = DecisionBoundaryDisplay(
        xx0=feat_1, xx1=feat_2, response=y[i],
        xlabel=iris.feature_names[0],
        ylabel=iris.feature_names[1],
)
    display.plot()
    display.ax_.scatter(
        iris.data[:, 0], iris.data[:, 1], c=iris.target, edgecolor="black")
    plt.title(f"k = {k_list[i]}")
```

