Kayvan Shah Assignment 2 solution

February 2, 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.

```
[3]: %%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: joblib>=1.1.1 in /usr/local/lib/python3.8/dist-packages (from scikit-learn) (1.2.0)
Requirement already satisfied: threadpoolctl>=2.0.0 in /usr/local/lib/python3.8/dist-packages (from scikit-learn) (3.1.0)

```
[4]: 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
[5]: iris = datasets.load iris()
     X = iris.data
     y = iris.target
[6]: 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)
[7]: X
[7]: 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],
            [5., 3.6, 1.4, 0.2],
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            [4.8, 3., 1.4, 0.1],
            [4.3, 3., 1.1, 0.1],
            [5.8, 4., 1.2, 0.2],
            [5.7, 4.4, 1.5, 0.4],
            [5.4, 3.9, 1.3, 0.4],
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```

```
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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, 3.8, 1.1],
[5.5, 2.4, 3.7, 1.],
[5.8, 2.7, 3.9, 1.2],
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[5.7, 2.5, 5., 2.],
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[6., 2.2, 5., 1.5],
[6.9, 3.2, 5.7, 2.3],
[5.6, 2.8, 4.9, 2.],
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[6.3, 2.7, 4.9, 1.8],
```

```
[6.7, 3.3, 5.7, 2.1],
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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.

```
[10]: ['DESCR',
       'data',
       'data_module',
       'feature_names',
       'filename',
       'frame',
       'target',
       'target_names']
[11]: iris.target_names
[11]: array(['setosa', 'versicolor', 'virginica'], dtype='<U10')</pre>
[12]: iris.feature_names
[12]: ['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
[13]: # Insert your answer here
      import pandas as pd
      iris_df = pd.DataFrame(iris.data, columns=iris.feature_names)
[14]: iris_df.describe()
[14]:
             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
                                                              1.000000
      min
                                          2.000000
      25%
                       5.100000
                                          2.800000
                                                              1.600000
      50%
                       5.800000
                                          3.000000
                                                              4.350000
      75%
                       6.400000
                                          3.300000
                                                              5.100000
      max
                       7.900000
                                          4.400000
                                                              6.900000
             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.

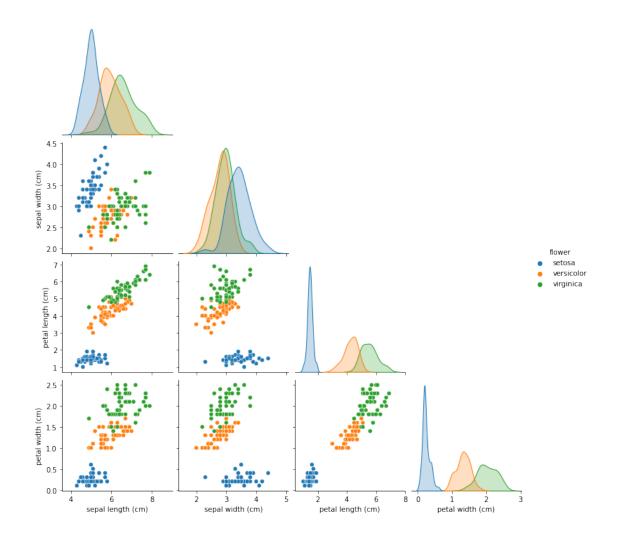
```
[15]:
         sepal length (cm) sepal width (cm) petal length (cm) petal width (cm)
                       5.1
                                         3.5
                                                            1.4
                                                                              0.2
                       4.9
      1
                                         3.0
                                                            1.4
                                                                              0.2
      2
                       4.7
                                         3.2
                                                            1.3
                                                                              0.2
      3
                       4.6
                                                            1.5
                                                                              0.2
                                         3.1
                       5.0
                                         3.6
                                                            1.4
                                                                              0.2
```

flower

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

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

[16]: <seaborn.axisgrid.PairGrid at 0x7feba20c1f40>



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.

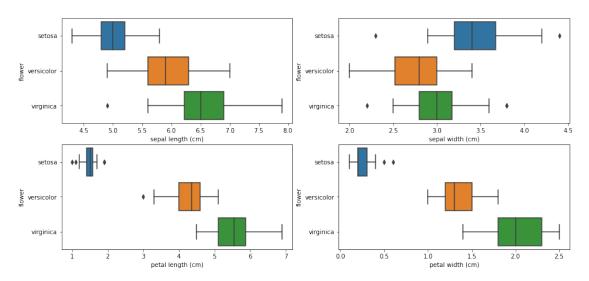
Some links to help you:

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

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

sns.boxplot(data=iris_df_pp, x=feat, y="flower", ax=axes[i])

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.htt + User guide https://scikit-learn.org/stable/modules/decomposition.html#pca

```
[18]: ### Insert your code here
from sklearn.decomposition import PCA

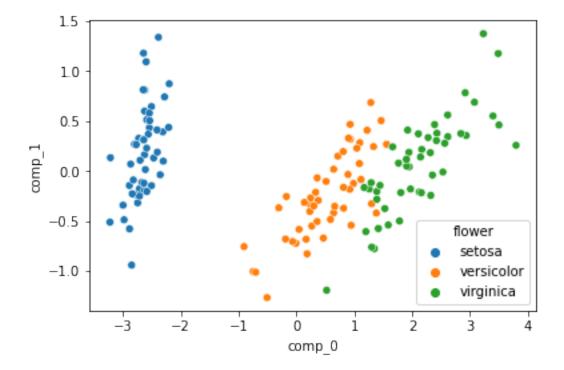
pca = PCA(n_components=2)
dimension_reducer = pca.fit(iris_df)
```

```
[18]: 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?

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

[19]: <matplotlib.axes._subplots.AxesSubplot at 0x7feb9f7622b0>



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.

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

```
[21]: 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.

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

model = KNeighborsClassifier(n_neighbors=5)
model.fit(X_train, y_train)
```

[22]: KNeighborsClassifier()

```
[23]: 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: 0.96 The test accuracy score is: 0.98

1.2.2 Part 2b Tuning hyperparameter k

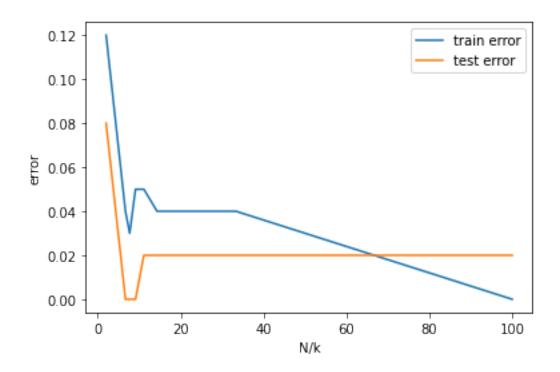
As we have seen in class, hyperparameter k of the K Nearest Neighbors classification affects the inductive bias. For this part train multiple near neighbor classifier models, store the results in a DataFrame. The plot plot training error and testing error versus N/k, where N=100.

1.2.3 Extension: Use different metric for knn classification.

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

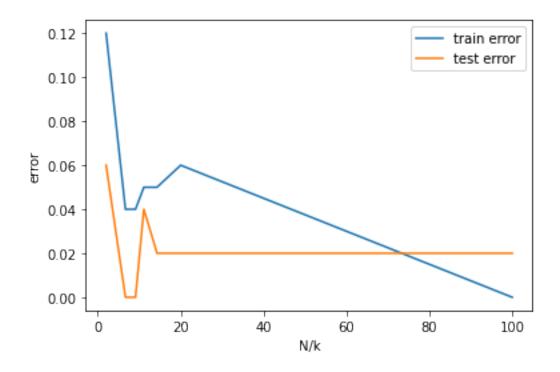
Distance Metrics Documentation: https://scikit-learn.org/stable/modules/generated/sklearn.metrics.pairwise.dist

```
[24]: k_list = [1, 3, 5, 7, 9, 11, 13, 15, 50]
      def get_train_test_error(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_train, y_train)
              train.append(1 - model.score(X_train, y_train))
              test.append(1 - 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
[25]: ### Insert your code
      # Use the `result` to store the DataFrame
      # euclidean
      result = get_train_test_error(metric="euclidean")
[26]: result.plot(x='N/k', y=['train error', 'test error'], ylabel='error')
```



```
[27]: ### Insert your code
# Use the `result` to store the DataFrame
# manhattan
result = get_train_test_error(metric="manhattan")
[28]: result.plot(x='N/k', y=['train error', 'test error'], ylabel='error')
```

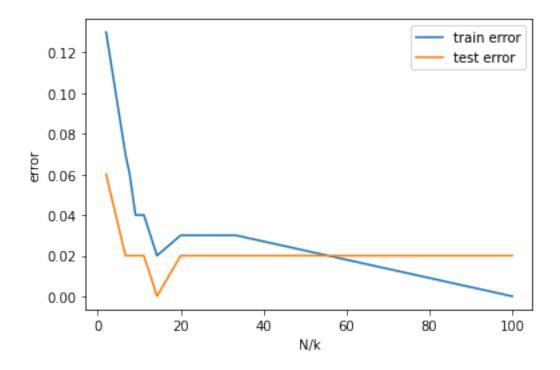
[28]: <matplotlib.axes._subplots.AxesSubplot at 0x7feb9a843100>



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

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

[30]: <matplotlib.axes._subplots.AxesSubplot at 0x7feb9a7bd310>



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

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

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

[2,0], [2,1], [2,2]]

```
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()
```

