

Solution_C

October 15, 2020

```
[1]: %matplotlib inline
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from sklearn.neighbors import KNeighborsClassifier
import filters, mutual_info, ttest
from sklearn.model_selection import train_test_split
from math import *
from sklearn.svm import SVC
from sklearn.metrics import accuracy_score, f1_score, confusion_matrix

[2]: data1 = pd.read_csv("Data/DLBCL.tsv",delimiter="\t", low_memory=False)
data1.drop(data1.index[[0,1]], inplace=True)
data1.dropna(inplace=True)

data2 = pd.read_csv("Data/leukemia.tsv",delimiter="\t", low_memory=False)
data2.drop(data2.index[[0,1]], inplace=True)
data2.dropna(inplace=True)

data3 = pd.read_csv("Data/lung.tsv",delimiter="\t", low_memory=False)
data3.drop(data3.index[[0,1]], inplace=True)
data3.dropna(inplace=True)
print("")

[3]: X1, y1 = data1.iloc[:, :-1], data1.iloc[:, -1]
X2, y2 = data2.iloc[:, 1:], data2.iloc[:, 0]
X3, y3 = data3.iloc[:, 1:], data3.iloc[:, 0]

[4]: X1_train, X1_test, y1_train, y1_test = train_test_split(X1, y1, test_size=0.25)
X2_train, X2_test, y2_train, y2_test = train_test_split(X2, y2, test_size=0.25)
X3_train, X3_test, y3_train, y3_test = train_test_split(X3, y3, test_size=0.25)

[5]: n1_features = len(X1.columns)
n2_features = len(X2.columns)
n3_features = len(X3.columns)
```

```
print("No. of features in DBCL data: {}".format(n1_features))
print("No. of features in Leukemia data: {}".format(n2_features))
print("No. of features in Lung data: {}".format(n3_features))
```

No. of features in DBCL data: 7070
 No. of features in Leukemia data: 5147
 No. of features in Lung data: 12600

```
[6]: n1_select = floor((0.2*n1_features)/3)
      n2_select = floor((0.2*n2_features)/3)
      n3_select = floor((0.2*n3_features)/3)

      print("No. of features to be selected from DBCL data (N/3): {}".
            ↳format(n1_select))
      print("No. of features to be selected from Leukemia data (N/3): {}".
            ↳format(n2_select))
      print("No. of features to be selected from Lung data (N/3): {}".
            ↳format(n3_select))
```

No. of features to be selected from DBCL data (N/3): 471
 No. of features to be selected from Leukemia data (N/3): 343
 No. of features to be selected from Lung data (N/3): 840

```
[7]: def topk_features_mi(X, y, n_features):
      fi = mutual_info.mutual_info_classif(X, y)
      f = X.columns
      fs = [j for i,j in sorted(zip(fi,f), reverse=True)][0:n_features]
      fi = [i for i,j in sorted(zip(fi,f), reverse=True)][0:n_features]
      Xs = X[fs]

      return Xs, y, fs, fi

      def topk_features_f_calssif(X, y, n_features):
          fi = filters.f_classif(X, y)
          fi = fi[0]
          f = X.columns
          fs = [j for i,j in sorted(zip(fi,f), reverse=True)][0:n_features]
          fi = [i for i,j in sorted(zip(fi,f), reverse=True)][0:n_features]
          Xs = X[fs]

          return Xs, y, fs, fi

      def topk_features_t_test(X, y, n_features):
          f,fi = ttest.get_features(X, y)
          #     fi = fi[0]
          #     print(f)
          #     f = X.columns
```

```

fs = [j for i,j in sorted(zip(fi,f), reverse=True)][[:n_features]
fi = [i for i,j in sorted(zip(fi,f), reverse=True)][[:n_features]
Xs = X[fs]

return Xs, y, fs, fi

def results_kNN(X_train, y_train, X_test, y_test):
    Clf = KNeighborsClassifier(n_neighbors=3)
    Clf.fit(X_train, y_train)

    y_pred = Clf.predict(X_test)
    acc = accuracy_score(y_test.values, y_pred)
    fscore = f1_score(y_test.values, y_pred, average='weighted')
    cnf_matrix = confusion_matrix(y_test, y_pred)

    return acc, fscore, cnf_matrix

def results_svm(X_train, y_train, X_test, y_test):
    Clf = SVC(kernel='rbf')
    Clf.fit(X_train, y_train)

    y_pred = Clf.predict(X_test)
    acc = accuracy_score(y_test, y_pred)
    fscore = f1_score(y_test, y_pred, average='weighted')
    cnf_matrix = confusion_matrix(y_test, y_pred)

    return acc, fscore, cnf_matrix

```

0.1 DBCL Dataset

0.1.1 F1 → F2 → F3

```

[8]: X1_train_mi, y1_train_mi, fs1_mi, fi1_mi = topk_features_mi(X1_train, y1_train,
    ↪3*n1_select)
X1_train_f, y1_train_f, fs1_f, fi1_f = topk_features_f_calssif(X1_train_mi,
    ↪y1_train_mi, 2*n1_select)
X1_train_t, y1_train_t, fs1_t, fi1_t = topk_features_t_test(X1_train_f,
    ↪y1_train_f, n1_select)

X1_test_t = X1_test[fs1_t]

```

/home/aquarius31/EndSem Notes/ML - Assignments/Assignments/Assignment
2/ttest.py:39: SettingWithCopyWarning:
A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row_indexer,col_indexer] = value instead

See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/user_guide/indexing.html#returning-a-view-versus-a-copy

```
df["class"] = pd.DataFrame(y)
```

```
[9]: knn_results1 = results_kNN(X1_train_t, y1_train_t, X1_test_t, y1_test)
      svm_results1 = results_svm(X1_train_t, y1_train_t, X1_test_t, y1_test)

      print("kNN Results for DBCL data:")
      print("Accuracy: {}".format(knn_results1[0]))
      print("Weighted F1-Score: {}".format(knn_results1[1]))
      print("Confusion Matrix:")
      print(knn_results1[2])
      print("\n")

      print("SVM Results for DBCL data:")
      print("Accuracy: {}".format(svm_results1[0]))
      print("Weighted F1-Score: {}".format(svm_results1[1]))
      print("Confusion Matrix:")
      print(svm_results1[2])
      print("\n")
```

```
kNN Results for DBCL data:
Accuracy: 0.9
Weighted F1-Score: 0.8933333333333333
Confusion Matrix:
[[14  0]
 [ 2  4]]
```

```
SVM Results for DBCL data:
Accuracy: 0.95
Weighted F1-Score: 0.94858934169279
Confusion Matrix:
[[14  0]
 [ 1  5]]
```

0.1.2 F2 → F3 → F1

```
[10]: X1_train_f, y1_train_f, fs1_f, fi1_f = topk_features_f_calssif(X1_train,
      ↪y1_train, n1_select)
      X1_train_t, y1_train_t, fs1_t, fi1_t = topk_features_t_test(X1_train_f,
      ↪y1_train_f, 2*n1_select)
      X1_train_mi, y1_train_mi, fs1_mi, fi1_mi = topk_features_mi(X1_train_t,
      ↪y1_train_t, n1_select)
```

```
X1_test_mi = X1_test[fs1_mi]
```

```
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2/ttest.py:39: SettingWithCopyWarning:  
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```
See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/user\_guide/indexing.html#returning-a-view-versus-a-copy  
df["class"] = pd.DataFrame(y)
```

```
[11]: knn_results1 = results_kNN(X1_train_mi, y1_train_mi, X1_test_mi, y1_test)  
      svm_results1 = results_svm(X1_train_mi, y1_train_mi, X1_test_mi, y1_test)  
  
      print("kNN Results for DBCL data:")  
      print("Accuracy: {}".format(knn_results1[0]))  
      print("Weighted F1-Score: {}".format(knn_results1[1]))  
      print("Confusion Matrix:")  
      print(knn_results1[2])  
      print("\n")  
  
      print("SVM Results for DBCL data:")  
      print("Accuracy: {}".format(svm_results1[0]))  
      print("Weighted F1-Score: {}".format(svm_results1[1]))  
      print("Confusion Matrix:")  
      print(svm_results1[2])  
      print("\n")
```

```
kNN Results for DBCL data:  
Accuracy: 0.95  
Weighted F1-Score: 0.94858934169279  
Confusion Matrix:  
[[14  0]  
 [ 1  5]]
```

```
SVM Results for DBCL data:  
Accuracy: 0.95  
Weighted F1-Score: 0.94858934169279  
Confusion Matrix:  
[[14  0]  
 [ 1  5]]
```

0.1.3 F3 → F1 → F2

```
[12]: X1_train_t, y1_train_t, fs1_t, fi1_t = topk_features_t_test(X1_train, y1_train,
    ↳ 3*n1_select)
X1_train_mi, y1_train_mi, fs1_mi, fi1_mi = topk_features_mi(X1_train_t,
    ↳ y1_train_t, 2*n1_select)
X1_train_f, y1_train_f, fs1_f, fi1_f = topk_features_f_calssif(X1_train_mi,
    ↳ y1_train_mi, n1_select)

X1_test_f = X1_test[fs1_f]
```

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See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/user_guide/indexing.html#returning-a-view-versus-a-copy
df["class"] = pd.DataFrame(y)

```
[13]: knn_results1 = results_kNN(X1_train_f, y1_train_f, X1_test_f, y1_test)
svm_results1 = results_svm(X1_train_f, y1_train_f, X1_test_f, y1_test)

print("kNN Results for DBCL data:")
print("Accuracy: {}".format(knn_results1[0]))
print("Weighted F1-Score: {}".format(knn_results1[1]))
print("Confusion Matrix:")
print(knn_results1[2])
print("\n")

print("SVM Results for DBCL data:")
print("Accuracy: {}".format(svm_results1[0]))
print("Weighted F1-Score: {}".format(svm_results1[1]))
print("Confusion Matrix:")
print(svm_results1[2])
print("\n")
```

kNN Results for DBCL data:
Accuracy: 0.95
Weighted F1-Score: 0.94858934169279
Confusion Matrix:
[[14 0]
 [1 5]]

SVM Results for DBCL data:

```

Accuracy: 0.95
Weighted F1-Score: 0.94858934169279
Confusion Matrix:
[[14  0]
 [ 1  5]]

```

[]:

0.2 Leukemia Dataset

0.2.1 F1 → F2 → F3

```

[14]: X2_train_mi, y2_train_mi, fs2_mi, fi2_mi = topk_features_mi(X2_train, y2_train,
    ↳ 3*n2_select)
X2_train_f, y2_train_f, fs2_f, fi2_f = topk_features_f_calssif(X2_train_mi,
    ↳ y2_train_mi, 2*n2_select)
X2_train_t, y2_train_t, fs2_t, fi2_t = topk_features_t_test(X2_train_f,
    ↳ y2_train_f, n2_select)

X2_test_t = X2_test[fs2_t]

```

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See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/user_guide/indexing.html#returning-a-view-versus-a-copy
df["class"] = pd.DataFrame(y)

```

[15]: knn_results2 = results_kNN(X2_train_t, y2_train_t, X2_test_t, y2_test)
svm_results2 = results_svm(X2_train_t, y2_train_t, X2_test_t, y2_test)

print("kNN Results for Leukemia data:")
print("Accuracy: {}".format(knn_results2[0]))
print("Weighted F1-Score: {}".format(knn_results2[1]))
print("Confusion Matrix:")
print(knn_results2[2])
print("\n")

print("SVM Results for Leukemia data:")
print("Accuracy: {}".format(svm_results2[0]))
print("Weighted F1-Score: {}".format(svm_results2[1]))
print("Confusion Matrix:")

```

```
print(svm_results2[2])
print("\n")
```

kNN Results for Leukemia data:

Accuracy: 1.0

Weighted F1-Score: 1.0

Confusion Matrix:

```
[[13  0]
 [ 0  5]]
```

SVM Results for Leukemia data:

Accuracy: 1.0

Weighted F1-Score: 1.0

Confusion Matrix:

```
[[13  0]
 [ 0  5]]
```

0.2.2 F2 → F3 → F1

```
[16]: X2_train_f, y2_train_f, fs2_f, fi2_f = topk_features_f_calssif(X2_train,
    ↪y2_train, n2_select)
X2_train_t, y2_train_t, fs2_t, fi2_t = topk_features_t_test(X2_train_f,
    ↪y2_train_f, 2*n2_select)
X2_train_mi, y2_train_mi, fs2_mi, fi2_mi = topk_features_mi(X2_train_t,
    ↪y2_train_t, n2_select)

X2_test_mi = X2_test[fs2_mi]
```

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2/ttest.py:39: SettingWithCopyWarning:

A value is trying to be set on a copy of a slice from a DataFrame.

Try using .loc[row_indexer,col_indexer] = value instead

See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/user_guide/indexing.html#returning-a-view-versus-a-copy

```
df["class"] = pd.DataFrame(y)
```

```
[17]: knn_results2 = results_kNN(X2_train_mi, y2_train_mi, X2_test_mi, y2_test)
svm_results2 = results_svm(X2_train_mi, y2_train_mi, X2_test_mi, y2_test)

print("kNN Results for Leukemia data:")
print("Accuracy: {}".format(knn_results2[0]))
print("Weighted F1-Score: {}".format(knn_results2[1]))
```



```

print("Confusion Matrix:")
print(knn_results2[2])
print("\n")

print("SVM Results for Leukemia data:")
print("Accuracy: {}".format(svm_results2[0]))
print("Weighted F1-Score: {}".format(svm_results2[1]))
print("Confusion Matrix:")
print(svm_results2[2])
print("\n")

```

kNN Results for Leukemia data:
Accuracy: 1.0
Weighted F1-Score: 1.0
Confusion Matrix:
[[13 0]
 [0 5]]

SVM Results for Leukemia data:
Accuracy: 1.0
Weighted F1-Score: 1.0
Confusion Matrix:
[[13 0]
 [0 5]]

0.2.3 F3 → F1 → F2

```

[18]: X2_train_t, y2_train_t, fs2_t, fi2_t = topk_features_t_test(X2_train, y2_train,
    ↳ 3*n2_select)
X2_train_mi, y2_train_mi, fs2_mi, fi2_mi = topk_features_mi(X2_train_t,
    ↳ y2_train_t, 2*n2_select)
X2_train_f, y2_train_f, fs2_f, fi2_f = topk_features_f_calssif(X2_train_mi,
    ↳ y2_train_mi, n2_select)

X2_test_f = X2_test[fs2_f]

```

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A value is trying to be set on a copy of a slice from a DataFrame.
Try using .loc[row_indexer,col_indexer] = value instead

See the caveats in the documentation: <http://pandas.pydata.org/pandas->

```
docs/stable/user_guide/indexing.html#returning-a-view-versus-a-copy
df["class"] = pd.DataFrame(y)
```

```
[19]: knn_results2 = results_kNN(X2_train_f, y2_train_f, X2_test_f, y2_test)
      svm_results2 = results_svm(X2_train_f, y2_train_f, X2_test_f, y2_test)

      print("kNN Results for Leukemia data:")
      print("Accuracy: {}".format(knn_results2[0]))
      print("Weighted F1-Score: {}".format(knn_results2[1]))
      print("Confusion Matrix:")
      print(knn_results2[2])
      print("\n")

      print("SVM Results for Leukemia data:")
      print("Accuracy: {}".format(svm_results2[0]))
      print("Weighted F1-Score: {}".format(svm_results2[1]))
      print("Confusion Matrix:")
      print(svm_results2[2])
      print("\n")
```

```
kNN Results for Leukemia data:
Accuracy: 1.0
Weighted F1-Score: 1.0
Confusion Matrix:
[[13  0]
 [ 0  5]]
```

```
SVM Results for Leukemia data:
Accuracy: 1.0
Weighted F1-Score: 1.0
Confusion Matrix:
[[13  0]
 [ 0  5]]
```

```
[ ]:
```

0.3 Lung Dataset

0.3.1 F1 → F2 → F3

```
[20]: X3_train_mi, y3_train_mi, fs3_mi, fi3_mi = topk_features_mi(X3_train, y3_train,
      ↪ 3*n3_select)
```

```

X3_train_f, y3_train_f, fs3_f, fi3_f = topk_features_f_calssif(X3_train_mi,
    ↪y3_train_mi, 2*n3_select)
X3_train_t, y3_train_t, fs3_t, fi3_t = topk_features_t_test(X3_train_f,
    ↪y3_train_f, n3_select)

X3_test_t = X3_test[fs3_t]

```

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See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/user_guide/indexing.html#returning-a-view-versus-a-copy
df["class"] = pd.DataFrame(y)

```

[21]: knn_results3 = results_kNN(X3_train_t, y3_train_t, X3_test_t, y3_test)
      svm_results3 = results_svm(X3_train_t, y3_train_t, X3_test_t, y3_test)

      print("kNN Results for Lung data:")
      print("Accuracy: {}".format(knn_results3[0]))
      print("Weighted F1-Score: {}".format(knn_results3[1]))
      print("Confusion Matrix:")
      print(knn_results3[2])
      print("\n")

      print("SVM Results for Lung data:")
      print("Accuracy: {}".format(svm_results3[0]))
      print("Weighted F1-Score: {}".format(svm_results3[1]))
      print("Confusion Matrix:")
      print(svm_results3[2])
      print("\n")

```

kNN Results for Lung data:
Accuracy: 0.8823529411764706
Weighted F1-Score: 0.8691736475497868
Confusion Matrix:
[[33 0 0 0 1]
 [0 2 0 0 0]
 [0 0 7 0 0]
 [1 0 0 1 1]
 [3 0 0 0 2]]

SVM Results for Lung data:
Accuracy: 0.8431372549019608
Weighted F1-Score: 0.8090580729899616

```
Confusion Matrix:
[[34  0  0  0  0]
 [ 0  2  0  0  0]
 [ 3  0  4  0  0]
 [ 3  0  0  0  0]
 [ 2  0  0  0  3]]
```

0.3.2 F2 → F3 → F1

```
[22]: X3_train_f, y3_train_f, fs3_f, fi3_f = topk_features_f_calssif(X3_train,
    ↪y3_train, n3_select)
X3_train_t, y3_train_t, fs3_t, fi3_t = topk_features_t_test(X3_train_f,
    ↪y3_train_f, 2*n3_select)
X3_train_mi, y3_train_mi, fs3_mi, fi3_mi = topk_features_mi(X3_train_t,
    ↪y3_train_t, n3_select)

X3_test_mi = X3_test[fs3_mi]
```

```
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```

```
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df["class"] = pd.DataFrame(y)
```

```
[23]: knn_results3 = results_kNN(X3_train_mi, y3_train_mi, X3_test_mi, y3_test)
svm_results3 = results_svm(X3_train_mi, y3_train_mi, X3_test_mi, y3_test)

print("kNN Results for Lung data:")
print("Accuracy: {}".format(knn_results3[0]))
print("Weighted F1-Score: {}".format(knn_results3[1]))
print("Confusion Matrix:")
print(knn_results3[2])
print("\n")

print("SVM Results for Lung data:")
print("Accuracy: {}".format(svm_results3[0]))
print("Weighted F1-Score: {}".format(svm_results3[1]))
print("Confusion Matrix:")
print(svm_results3[2])
print("\n")
```

kNN Results for Lung data:

```

Accuracy: 0.9607843137254902
Weighted F1-Score: 0.9597147950089127
Confusion Matrix:
[[33  0  0  0  1]
 [ 0  2  0  0  0]
 [ 0  0  7  0  0]
 [ 1  0  0  2  0]
 [ 0  0  0  0  5]]

```

```

SVM Results for Lung data:
Accuracy: 0.9215686274509803
Weighted F1-Score: 0.8941685765215177
Confusion Matrix:
[[33  0  0  0  1]
 [ 0  2  0  0  0]
 [ 0  0  7  0  0]
 [ 3  0  0  0  0]
 [ 0  0  0  0  5]]

```

0.3.3 F3 → F1 → F2

```

[24]: X3_train_t, y3_train_t, fs3_t, fi3_t = topk_features_t_test(X3_train, y3_train,
    ↳ 3*n3_select)
X3_train_mi, y3_train_mi, fs3_mi, fi3_mi = topk_features_mi(X3_train_t,
    ↳ y3_train_t, 2*n3_select)
X3_train_f, y3_train_f, fs3_f, fi3_f = topk_features_f_calssif(X3_train_mi,
    ↳ y3_train_mi, n3_select)

X3_test_f = X3_test[fs3_f]

```

```

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```

```

See the caveats in the documentation: http://pandas.pydata.org/pandas-docs/stable/user\_guide/indexing.html#returning-a-view-versus-a-copy
df["class"] = pd.DataFrame(y)

```

```

[25]: knn_results3 = results_kNN(X3_train_f, y3_train_f, X3_test_f, y3_test)
svm_results3 = results_svm(X3_train_f, y3_train_f, X3_test_f, y3_test)

```

```

print("kNN Results for Lung data:")
print("Accuracy: {}".format(knn_results3[0]))
print("Weighted F1-Score: {}".format(knn_results3[1]))
print("Confusion Matrix:")
print(knn_results3[2])
print("\n")

print("SVM Results for Lung data:")
print("Accuracy: {}".format(svm_results3[0]))
print("Weighted F1-Score: {}".format(svm_results3[1]))
print("Confusion Matrix:")
print(svm_results3[2])
print("\n")

```

kNN Results for Lung data:
 Accuracy: 0.9215686274509803
 Weighted F1-Score: 0.9128851540616246
 Confusion Matrix:
 [[33 0 0 0 1]
 [0 2 0 0 0]
 [0 0 7 0 0]
 [2 0 0 1 0]
 [1 0 0 0 4]]

SVM Results for Lung data:
 Accuracy: 0.8431372549019608
 Weighted F1-Score: 0.8120634002986944
 Confusion Matrix:
 [[33 0 0 0 1]
 [0 2 0 0 0]
 [3 0 4 0 0]
 [3 0 0 0 0]
 [1 0 0 0 4]]

[]:

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