General

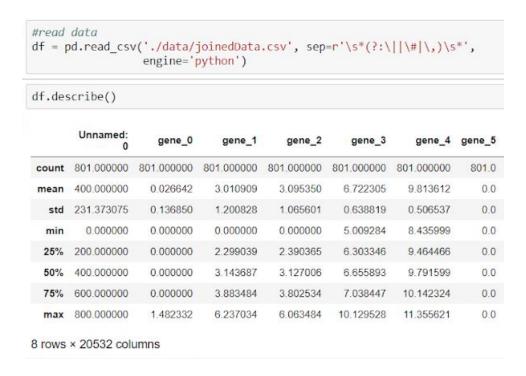
The current prototype that is working are all three classifiers, the KNN, DNN and OVR. The author had used tensorflow and sklearn website and libraries to achieve this, apart from numpy, pandas etc. She used the sample codes that was provided by the websites tensorflow and sklearn and changed it a bit to suit the authors needs. The reading in, standardizing and splitting the data sets are the authors work. She will then try to use less built in features and create her own code.

One-Vs-Rest

```
In [2]: #read the data set
        df = pd.read_csv('./data/joinedData.csv', sep=r'\s*(?:\||\#|\,)\s*',
           engine='python')
In [3]: from sklearn.feature_selection import RFE
        #change the 5 tumour types to numbers
        Class = {'LUAD': 0, 'BRCA': 1, 'KIRC': 2, 'PRAD': 3, 'COAD': 4}
        #this is where we add the class to the table
        df.Class = [Class[item] for item in df.Class]
        #drop the 2 unnamed table because we do not need them
        df = df.drop('Unnamed: 0',1)
       df = df.drop('Unnamed: 0.1',1)
       df
       4 1 0.000000 2.655741 2.821547 6.539454 9.738265 0.0 6.566967 0.36 A
         5
             3 0.000000 3.467853 3.581918 6.620243 9.706829 0.0 7.758510 0.00
        6 2 0.000000 1.224966 1.691177 6.572007 9.640511 0.0 6.754888 0.53
             3 0.000000 2.854853 1.750478 7.226720 9.758691
                                                             0.0 5.952103 0.00
        8 1 0.000000 3.992125 2.772730 6.546692 10.488252 0.0 7.690222 0.35
                                                             0.0 7.947216 0.72
             3 0.000000 3.642494 4.423558 6.849511 9.464466
         10 1 0.000000 3.492071 3.553373 7.151707 10.253446 0.0 8.301258 0.00
         11
               2 0.000000 2.941181 2.663276 6.561690 9.376293
                                                             0.0 7.860323 0.75
               3 0.000000 3.970348 2.364292 7.145443 9.240605
         12
                                                             0.0 7.810758 0.00
                1 0.000000 1.551048 3.529846 6.326825 10.633849
         14
               1 0.000000 1.964842 2.183010 6.596832 10.248141
                                                             0.0 7.087251 0.44
         15
             1 0.000000 2.901379 3.685368 6.669665 9.999098
                                                             0.0 6.948834 0.00
        16 0 0.000000 3.460913 3.618474 5.661048 9.731217 0.0 8.435591 1.03 .
                                                              >
In [4]: X = df.drop('Class', axis=1).values
       y = df['Class'].values
       y = np.asarray(y)
```

```
n [6]: #Standardize data
       X = (X - X.mean()) / (X.max() - X.min())
       #Split the training and testing data
       X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.20
       X_train
       4
ut[6]: array([[-0.31009073, -0.17314025, -0.17092733, ..., 0.14479179,
               -0.16756171, -0.31009073],
[-0.31009073, -0.24838072, -0.1657764 , ..., 0.15107929,
                -0.02477067, -0.31009073],
               [-0.31009073, -0.08768639, -0.12152632, ..., 0.15792214,
                0.0799575 , -0.31009073],
               [-0.31009073, -0.20680766, -0.14623389, ..., 0.14587579,
                -0.03078254, -0.31009073],
               [-0.31009073, -0.11563346, -0.12269672, ..., 0.15631198,
               -0.14153606, -0.29298897],
[-0.31009073, -0.19012855, -0.15522276, ..., 0.15125456,
                -0.09566869, -0.31009073]])
n [7]: ##Predict
       df2 = df.drop('Class',1)
       test = df2.iloc[[800]]
       test
ut[7]:
             gene_0 gene_1 gene_2 gene_3 gene_4 gene_5 gene_6 gene_7 gene_8
        800 0.0 2.325242 3.805932 6.530246 9.560367
                                                         0.0 7.957027
       1 rows × 20531 columns
       4
```

K-Nearest Neighbour



```
In [4]: #change the 5 tumour types to numbers
        Class = {'LUAD': 0, 'BRCA': 1, 'KIRC': 2, 'PRAD': 3, 'COAD': 4}
        #this is where we add the class to the table
        df.Class = [Class[item] for item in df.Class]
        #drop the 2 unnamed table because we do not need them
        df = df.drop('Unnamed: 0',1)
        df = df.drop('Unnamed: 0.1',1)
        #Split the X and y
        X = df.drop('Class', axis=1).values
        y = df['Class'].values
        y = np.asarray(y)
        #Standarize using min and max
        X = (X - X.mean()) / (X.max() - X.min())
        #Split the data set with 80 to traina dn20 to test
        X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.20
        X_train
```

Neural Networks

```
#read the data
df = pd.read_csv('./data/joinedData.csv', sep=r'\s*(?:\|\#|\,)\s*',
                 engine='python')
#change the classes to numbers
Class = {'LUAD': 0, 'BRCA': 1, 'KIRC': 2, 'PRAD': 3, 'COAD': 4}
df.Class = [Class[item] for item in df.Class]
df = df.drop('Unnamed: 0',1)
df = df.drop('Unnamed: 0.1',1)
df
#separate the data into X and y
X = df.drop('Class', axis=1).values
y = df['Class'].values
y = np.asarray(y)
#standardize the data
X = (X - X.mean()) / (X.max() - X.min())
#Split the data set to test and train
X train, X test, y train, y test = train test split(X, y, test size = 0.20
X train
```

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