SVC Assignment

Part 1 Dos vs Non Dos

```
In [*]: import numpy as np
    import pandas as pd
    import matplotlib.pyplot as plt
    import time
    from sklearn.linear_model import LogisticRegression, Lasso, Ridge
    from sklearn.preprocessing import OneHotEncoder
    from sklearn.compose import make_column_transformer
    from sklearn.preprocessing import scale
    from sklearn.model_selection import train_test_split
    from sklearn import svm, datasets
    from sklearn import metrics
    from sklearn.svm import SVC
    from sklearn.preprocessing import MinMaxScaler
```

```
In [74]: df = pd.read_csv('kddcup99_csv.csv')
```

In [75]: df

| | duration | protocol_type | service | flag | src_bytes | dst_bytes | land | wrong_fragment | urgent |
|--------|----------|---------------|---------|------|-----------|-----------|------|----------------|--------|
| 0 | 0 | tcp | http | SF | 181 | 5450 | 0 | 0 | 0 |
| 1 | 0 | tcp | http | SF | 239 | 486 | 0 | 0 | 0 |
| 2 | 0 | tcp | http | SF | 235 | 1337 | 0 | 0 | 0 |
| 3 | 0 | tcp | http | SF | 219 | 1337 | 0 | 0 | 0 |
| 4 | 0 | tcp | http | SF | 217 | 2032 | 0 | 0 | 0 |
| | *** | | | | | | | | ••• |
| 494015 | 0 | tcp | http | SF | 310 | 1881 | 0 | 0 | 0 |
| 494016 | 0 | tcp | http | SF | 282 | 2286 | 0 | 0 | 0 |
| 494017 | 0 | tcp | http | SF | 203 | 1200 | 0 | 0 | 0 |
| 494018 | 0 | tcp | http | SF | 291 | 1200 | 0 | 0 | 0 |
| 494019 | 0 | tcp | http | SF | 219 | 1234 | 0 | 0 | 0 |

```
In [4]: y = df.iloc[0:,4:40].values
#X = df.iloc[0:,4:40].values
#probe = ['ipsweep.','satan.','nmap.','portsweep.']
y = np.where(((df['label']== 'back') | (df['label'] == 'land') | (df['label'] == 'np.unique(y))
r
```

Out[4]: array([0, 1])

Part 2: Running SVM model four times with different kernels

```
In [9]:
         Xs = MinMaxScaler().fit transform(X1)
         from imblearn.under sampling import RandomUnderSampler
         ros = RandomUnderSampler(random state=0)
         X underresampled, y underresampled = ros.fit resample (Xs,y)
         from sklearn.model selection import train_test_split
         X_train, X_test, y_train, y_test = train_test_split(X_underresampled,y_unde
         X train
 Out[9]: array([[0., 1., 0., ..., 0., 0., 0.],
                [1., 0., 0., ..., 0., 0., 0.]
                [0., 1., 0., ..., 1., 0., 0.],
                [0., 0., 1., ..., 0., 0., 0.],
                [1., 0., 0., ..., 0., 0., 0.]
                [1., 0., 0., ..., 0., 0., 0.]]
 In [ ]: | svc = svm.SVC(probability=False, kernel="linear", C=2.8, gamma=.0073,verbo
         svc.fit(X train, y train)
         svc.score(X test,y test)
         [LibSVM]
In [12]: svc.score(X_train,y_train)
Out[12]: 0.9994515505883643
```

```
In [13]: svc = svm.SVC(probability=False, kernel="rbf", C=2.8, gamma=.0073,verbose=
         svc.fit(X train, y train)
         svc.score(X_test,y_test)
         [LibSVM]
Out[13]: 0.9958074344911639
In [14]: svc.score(X_train,y_train)
Out[14]: 0.9965691442361013
In [15]: svc = svm.SVC(probability=False, kernel="sigmoid", C=2.8, gamma=.0073,verb
         svc.fit(X train, y train)
         svc.score(X_test,y_test)
         [LibSVM]
Out[15]: 0.9957586837294333
In [16]: svc.score(X_train,y_train)
Out[16]: 0.9964533604714227
In [17]: svc = svm.SVC(probability=False, kernel="poly", C=2.8, gamma=.0073, verbose
         svc.fit(X_train, y_train)
         svc.score(X test,y test)
         [LibSVM]
Out[17]: 0.9926630103595369
In [18]: | svc.score(X train, y train)
Out[18]: 0.9931565701192573
```

Compare the results for each of the kernels. Discuss the pros and cons of using each of the kernels that you've chosen.

Linear: Score test: 0.9992931139549055 Linear: Score train: 0.9994515505883643

RBF: Score test: 0.9994515505883643 Score train: 0.9965691442361013

Sigmoid: Score test: 0.9957586837294333 Score train: 0.9964533604714227

Poly: Score test: 0.9926630103595369 train test: 0.9931565701192573

The cons of using all these kernels is that they all took a very long time to load and show the outputs. Poly took the longest out of all but had the lowest numbers. Linear had the highest numbers compared to all of them. Sigmoids score test and train test were close together. RBF had numbers very apart from each other with the score test and score train.

Part 3: Pick two features

```
In [77]: X=df.iloc[0:, [5, 23]].values
Out[77]: array([[5450,
                          8],
                [ 486,
                          8],
                          8],
                [1337,
                [1200,
                         18],
                         12],
                [1200,
                [1234,
                         3511)
In [78]: Xs = MinMaxScaler().fit transform(X)
         from imblearn.under sampling import RandomUnderSampler
         ros = RandomUnderSampler(random state=0)
         X_underresampled, y_underresampled = ros.fit_resample (Xs,y)
In [79]: from sklearn.model selection import train test split
         X_train, X_test, y_train, y_test = train_test_split(X_underresampled,y_unde
         X train
Out[79]: array([[0.00000000e+00, 1.00000000e+00],
                [1.72826211e-04, 1.76125245e-02],
                [4.74641681e-04, 1.95694716e-03],
                [0.00000000e+00, 1.0000000e+00],
                [2.71168398e-04, 5.87084149e-03],
                [5.12077662e-05, 5.28375734e-02]])
 In [*]: svc = svm.SVC(probability=False, kernel="rbf", C=2.8, gamma=.0073,verbose=
         svc.fit(X train, y train)
         svc.score(X_test,y_test)
```

[LibSVM]

```
In [*]: from matplotlib.colors import ListedColormap
        def plot decision regions(X, y, classifier, resolution=0.02):
            # setup marker generator and color map
            markers = ('s', 'x', 'o', '^i, 'v')
            colors = ('red', 'blue', 'lightgreen', 'gray', 'cyan')
            cmap = ListedColormap(colors[:len(np.unique(y))])
            # plot the decision surface
            x1_{\min}, x1_{\max} = X[:, 0].min() - 1, X[:, 0].max() + 1
            x2_{min}, x2_{max} = X[:, 1].min() - 1, X[:, 1].max() + 1
            xx1, xx2 = np.meshgrid(np.arange(x1 min, x1 max, resolution),
                                    np.arange(x2_min, x2_max, resolution))
            Z = classifier.predict(np.array([xx1.ravel(), xx2.ravel()]).T)
            Z = Z.reshape(xx1.shape)
            plt.contourf(xx1, xx2, Z, alpha=0.3, cmap=cmap)
            plt.xlim(xx1.min(), xx1.max())
            plt.ylim(xx2.min(), xx2.max())
            # plot class examples
            for idx, cl in enumerate(np.unique(y)):
                plt.scatter(x=X[y == cl, 0],
                            y=X[y == c1, 1],
                             alpha=0.8,
                             c=colors[idx],
                             marker=markers[idx],
                             label=cl,
                             edgecolor='black')
In [*]: plot_decision_regions(X_train, y_train, classifier=svc)
        plt.xlabel('sepal length [cm]')
        plt.ylabel('petal length [cm]')
        plt.legend(loc='upper left')
```

```
# plt.savefig('images/02 08.png', dpi=300)
plt.show()
```

```
In [*]: | svc = svm.SVC(probability=False, kernel="linear", C=2.8, gamma=.0073,verbo
        svc.fit(X train, y train)
        svc.score(X test,y test)
```

```
In [*]: from matplotlib.colors import ListedColormap
        def plot decision regions(X, y, classifier, resolution=0.02):
            # setup marker generator and color map
            markers = ('s', 'x', 'o', '^i, 'v')
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            cmap = ListedColormap(colors[:len(np.unique(y))])
            # plot the decision surface
            x1_{min}, x1_{max} = X[:, 0].min() - 1, X[:, 0].max() + 1
            x2_{min}, x2_{max} = X[:, 1].min() - 1, X[:, 1].max() + 1
            xx1, xx2 = np.meshgrid(np.arange(x1 min, x1 max, resolution),
                                    np.arange(x2_min, x2_max, resolution))
            Z = classifier.predict(np.array([xx1.ravel(), xx2.ravel()]).T)
            Z = Z.reshape(xx1.shape)
            plt.contourf(xx1, xx2, Z, alpha=0.3, cmap=cmap)
            plt.xlim(xx1.min(), xx1.max())
            plt.ylim(xx2.min(), xx2.max())
            # plot class examples
            for idx, cl in enumerate(np.unique(y)):
                plt.scatter(x=X[y == cl, 0],
                            y=X[y == c1, 1],
                             alpha=0.8,
                             c=colors[idx],
                             marker=markers[idx],
                             label=cl,
                             edgecolor='black')
```

```
In [*]: plot_decision_regions(X_test,y_test, classifier=svc)
    plt.xlabel('sepal length [cm]')
    plt.ylabel('petal length [cm]')
    plt.legend(loc='upper left')

# plt.savefig('images/02_08.png', dpi=300)
    plt.show()
```

Discuss your observations.

I saw that the linear feature has a higher percentage than the other feature that is RBF. The decision boundaries however looked very similar between these two.