```
In [1]: import collections
        import itertools
        import numpy as np
        import pandas as pd
        import scipy.spatial as sp
        import scipy.cluster.hierarchy as hc
        import matplotlib.pyplot as plt
        plt.style.use('ggplot')
        import matplotlib.colors as colors
        import matplotlib.cm as cmx
        import seaborn as sns
        from sklearn.preprocessing import StandardScaler
        from sklearn.decomposition import PCA
        from sklearn.linear_model import LogisticRegression
        from sklearn import svm
        from sklearn.model selection import GridSearchCV, train test split
        from sklearn.metrics import accuracy score, confusion matrix, classificatio
```

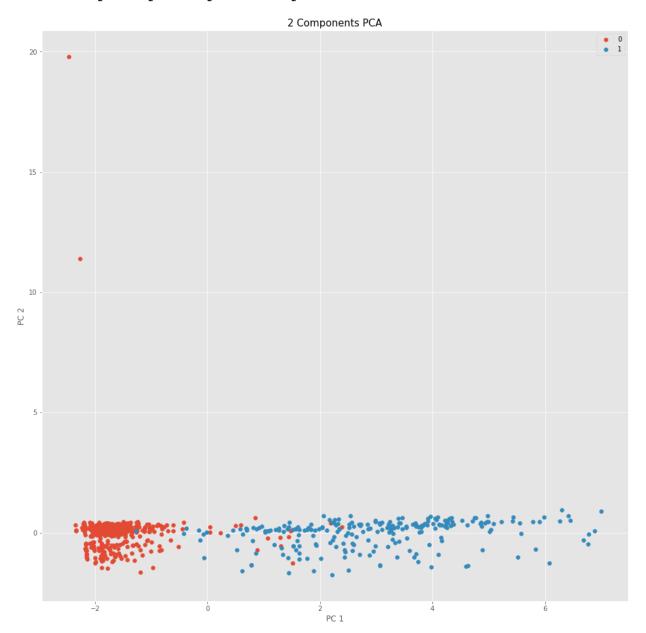
```
In [2]: colors(categorical):
        list(set(categorical))
       rical_to_number = pd.factorize(categorical)[0]
        plt.get cmap('brg') # https://matplotlib.org/tutorials/colors/colormaps.htm
        = colors.Normalize(vmin=0, vmax=len(uniq))
        Map = cmx.ScalarMappable(norm=cnorm, cmap=cmap)
        map = [scalarMap.to_rgba(i) for i in categorical_to_number]
        color map
       m pca(X, y, perc, visualize=True):
       ls = StandardScaler().fit_transform(X)
       PCA(perc)
       pal_comps = pca.fit_transform(std_vals)
       pal df = pd.DataFrame(data=principal comps)
        'Number of principal components explain {}% variance of data is {}'.format(r
       ualize:
       g = plt.figure(figsize=(16, 16))
        = fig.add_subplot(1, 1, 1)
       .set xlabel('PC 1', fontsize=12)
       .set ylabel('PC 2', fontsize=12)
       .set title('2 Components PCA', fontsize=15)
       r group in set(y):
         idx = np.where(y == group)[0]
         ax.scatter(principal df.loc[idx, 0], principal df.loc[idx, 1], label=group
       .legend()
       .grid(True)
       t.show()
       t(pca.explained variance ratio )
       t(principal df.values)
       rn pca.explained variance ratio , principal df.values
        None
       p(X, y, row_linkage, col_linkage):
       s.clustermap(
        figsize=(16,16), z_score=0,
        w linkage=feature linkage, col linkage=sample linkage,
       l colors=assign colors(y)
       DW()
       ibsvm format(url, classA, classB):
```

```
d.read_csv(url, sep=' ', header=None)
f.dropna(axis=1, how='all') # drop empty columns
f.dropna() #drop all rows that have any NaN values
= df.iloc[:,0].values
[np.where(labels==classA)] = 0
[np.where(labels==classB)] = 1
f.iloc[:,1:].applymap(lambda x: x.split(':')[1])
ass' | = labels
f.astype(float)
ass'] = df['class'].astype(int)
onfusion matrix(cm, classes, normalize=False, title='Confusion matrix', cmar
malize:
= cm.astype('float') / cm.sum(axis=1)[:, np.newaxis]
show(cm, interpolation='nearest', cmap=cmap)
tle(title)
lorbar()
arks = np.arange(len(classes))
icks(tick marks, classes, rotation=45)
icks(tick_marks, classes)
'.2f' if normalize else 'd'
= cm.max() / 2.
j in itertools.product(range(cm.shape[0]), range(cm.shape[1])):
t.text(j, i, format(cm[i, j], fmt),
      horizontalalignment='center',
       color='white' if cm[i, j] > thresh else 'black')
abel('True label')
abel('Predicted label')
ght layout()
(clf, model):
esult = clf.best score
Best classification accuracy of {} models: {}'.format(model, best result))
arameters = clf.best params
'Parameters of the best {} model: {}'.format(model, best parameters))
fy(X, y, classifier, grid params, name):
GridSearchCV(classifier, grid params, scoring='accuracy',
             cv=5, verbose=0, n jobs=-1) # for detailed log verbose=2
t(X, y)
(clf, name)
```

```
In [3]: breast = 'https://www.csie.ntu.edu.tw/~cjlin/libsvmtools/datasets/binary/br
    df = read_libsvm_format(breast, 2, 4)
    X = df.iloc[:,:-1].values
    y = df.iloc[:,-1].values
```

In [4]: perform\_pca(X, y, 0.90)

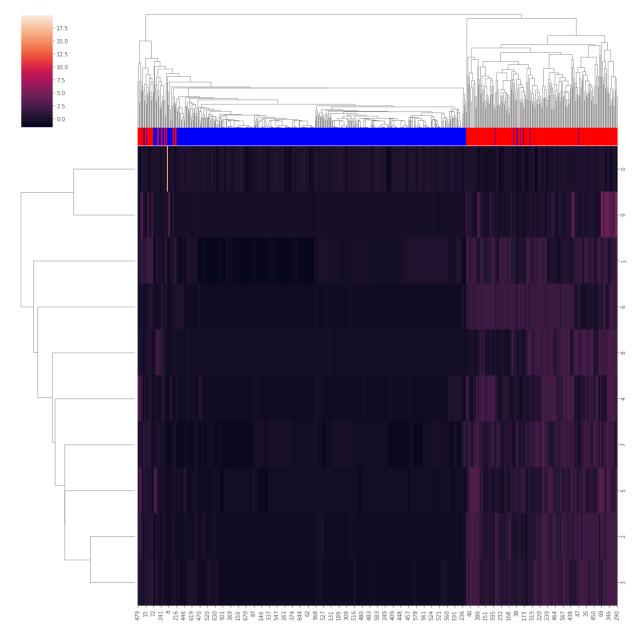
Number of principal components explain 90.0% variance of data is 6



```
In [5]: sample_dist = sp.distance.pdist(X, 'euclidean')
    feature_dist = sp.distance.pdist(X.T, 'euclidean')

sample_linkage = hc.linkage(sample_dist, method='average')
    feature_linkage = hc.linkage(feature_dist, method='average')

heatmap(X.T, y, sample_linkage, feature_linkage)
```



Best classification accuracy of SVM models: 0.9736904250751396

Parameters of the best SVM model: {'C': 1, 'class\_weight': 'balanced', 'd egree': 3, 'kernel': 'poly'}

Best classification accuracy of Logistic Regression models: 0.97077071704 59425

Parameters of the best Logistic Regression model: {'C': 100, 'class\_weigh t': 'balanced', 'penalty': 'l1', 'solver': 'saga'}

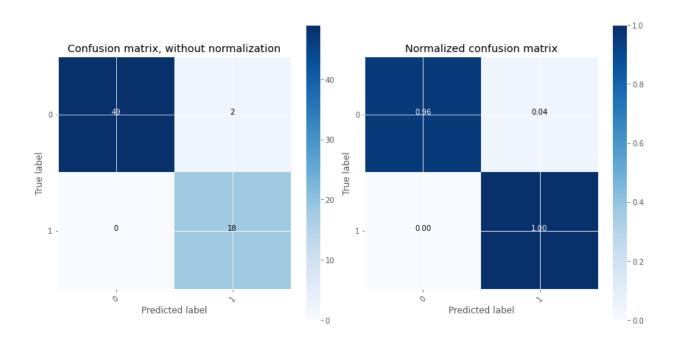
/Users/aryantimla/opt/anaconda3/lib/python3.8/site-packages/sklearn/linea r\_model/\_sag.py:328: ConvergenceWarning: The max\_iter was reached which m eans the coef did not converge

warnings.warn("The max iter was reached which means "

```
In [8]: clf = svm.SVC(gamma='scale', C=1, class_weight=None, degree=2, kernel='poly
        X train, X test, y train, y test = train_test_split(X, y, test_size=0.1, ra
        clf.fit(X_train, y_train)
        y_pred = clf.predict(X_test)
        print('Accuracy: {}\n'.format(accuracy score(y test, y pred)))
        print(classification_report(y_test, y_pred, target_names=[str(i) for i in s
        cnf matrix = confusion_matrix(y_test, y_pred)
        plt.subplots(figsize=(12,6))
        plt.subplot(1, 2, 1)
        plot confusion matrix(cnf matrix, classes=set(y test),
                              title='Confusion matrix, without normalization')
        plt.subplot(1, 2, 2)
        plot confusion matrix(cnf matrix, classes=set(y test), normalize=True,
                              title='Normalized confusion matrix')
        plt.tight layout()
        plt.show()
```

Accuracy: 0.9710144927536232

	precision	recall	f1-score	support
0	1.00	0.96	0.98	51
1	0.90	1.00	0.95	18
accuracy			0.97	69
macro avg	0.95	0.98	0.96	69
weighted avg	0.97	0.97	0.97	69



In [ ]: