# ECE 657A - Assignment 2

Date Submitted: Mar 09, 2022

#### **Importing Libraries**

```
In [1]:
       import numpy as np
       import pandas as pd
       import random
       from pprint import pprint
       from functools import reduce
       from scipy import stats
       from sklearn import neighbors
       from sklearn.preprocessing import MinMaxScaler
       from sklearn.model selection import train test split
       from sklearn.neighbors import KNeighborsClassifier
       from sklearn import metrics
       from sklearn.metrics import
       accuracy score, confusion matrix, ConfusionMatrixDisplay, classification report, matrix
        precision score, recall score, f1 score
       from sklearn.decomposition import PCA
       from sklearn.discriminant analysis import LinearDiscriminantAnalysis
       from sklearn.manifold import TSNE
       from sklearn.model selection import
       cross val score, KFold, StratifiedKFold, GridSearchCV, cross validate
       from sklearn.naive bayes import MultinomialNB, ComplementNB, GaussianNB
       from sklearn.tree import DecisionTreeClassifier
       from sklearn import tree
       import graphviz
       from sklearn.ensemble import
       RandomForestClassifier, GradientBoostingClassifier, AdaBoostClassifier
       import matplotlib.pyplot as plt
       import seaborn as sns
       sns.set(style="ticks", color codes=True)
       import warnings
       warnings.filterwarnings("ignore")
```

## Wine Dataset

## Preprocessing

## **Loading Data**

```
In [2]:
      D = ['fixed acidity', 'volatile acidity', 'citric acid', 'residual sugar',
       'chlorides', 'free sulfur dioxide', 'total sulfur dioxide', 'density', 'pH',
        'sulphates', 'alcohol']
       L = 'quality'
       C = 'color'
       DL = D + [L]
       DC = D + [C]
       DLC = DL + [C]
       #Loading Data set
       wine r = pd.read csv("winequality-red.csv", sep=';')
       #Loading Data set
       wine w = pd.read csv("winequality-white.csv", sep=';')
       wine w= wine w.copy()
       wine w[C] = np.zeros(wine w.shape[0])
       wine r[C] = np.ones(wine r.shape[0])
       wine = pd.concat([wine w, wine r])
       wine['color'] = wine['color'].astype('category')
       wine['quality'] = wine['quality'].astype('category')
```

### Normalization and Scaling

```
In [3]: #Z-Score Normalization
  wine_z = wine.copy()
  wine_z = wine_z.drop(columns = ['color' , 'quality'])
  wine_znormalized = wine_z.apply(stats.zscore)
  wine_znormalized['quality'] = wine['quality']
  wine_znormalized['color'] = wine['color']

  target_z = wine_znormalized['quality']
  data_z = wine_znormalized.drop(columns = 'quality')

  X_train_z, X_test_z, Y_train_z, Y_test_z = train_test_split(data_z, target_z, test_size=0.2 , random_state=27)
```

```
In [4]: #MinMAx Scaling
  wine_s = wine.copy()
  wine_s = wine_s.drop(columns = ['color' , 'quality'])
  wine_minmax = (wine_s - wine_s.min()) / ( wine_s.max() - wine_s.min())
  wine_minmax['quality'] = wine['quality']
  wine_minmax['color'] = wine['color']

wine_min_max = wine_minmax
  target = wine_min_max['quality']
```

```
data = wine_min_max.drop(columns = 'quality')
X_train, X_test, Y_train, Y_test = train_test_split(data, target,
test_size=0.2 , random_state=27)
```

### **Best formulated KNN from Assignment-1**

```
In [5]: knn = KNeighborsClassifier(n_neighbors=23 , metric = 'manhattan' , weights =
    "distance")
    knn.fit(X_train, Y_train)
    Y_pred = knn.predict(X_test)

    knn_train_acc = knn.score(X_train, Y_train)
    print("Training Score: ", knn_train_acc)
    knn_test_acc = knn.score(X_test, Y_test)
    print("Testing Score: ", knn_test_acc)
```

Training Score: 1.0 Testing Score: 0.68

## **Representation Learning**

#### Independent and dependent variable

```
In [6]: #Scaled 'data' and 'target'
    x=data
    y=target
    n_components=2
    target_names = np.sort(y.unique())
```

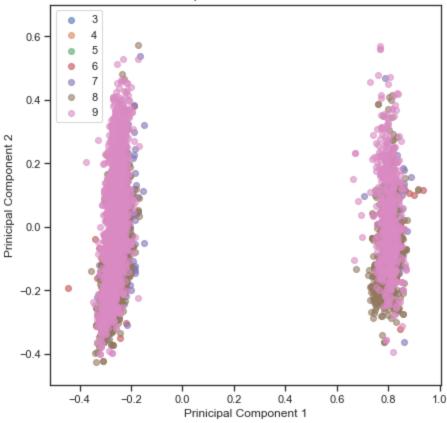
#### PCA

```
plt.ylim(-0.5,0.7)
plt.legend(loc="best", shadow=False, scatterpoints=1)
plt.xlabel("Prinicipal Component 1")
plt.ylabel("Prinicipal Component 2")
plt.title("First Two Components from PCA of wine dataset")
#plt.set_cmap("hsv")

plt.show()
```

```
PCA Model - Explained Variance of each component
Explained Variance Ratio
0 0.692196
1 0.118142
```

#### First Two Components from PCA of wine dataset

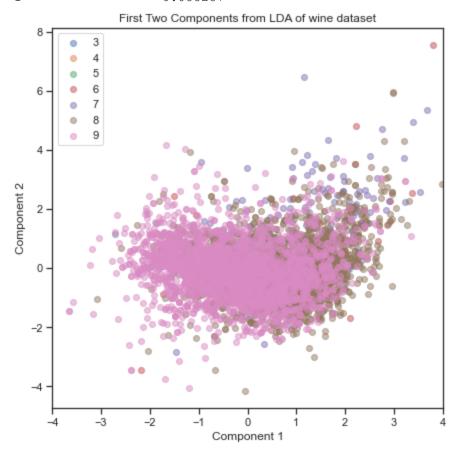


#### LDA

```
X_r2[y == i, 0], X_r2[y == i, 1], alpha=0.5, lw=1, label=target_name,
cmap="twilight"
)
plt.xlim(-4,4)
plt.legend(loc="best", shadow=False, scatterpoints=1)
plt.xlabel("Component 1")
plt.ylabel("Component 2")
plt.title("First Two Components from LDA of wine dataset")
#plt.set_cmap("twilight")

plt.show()
```

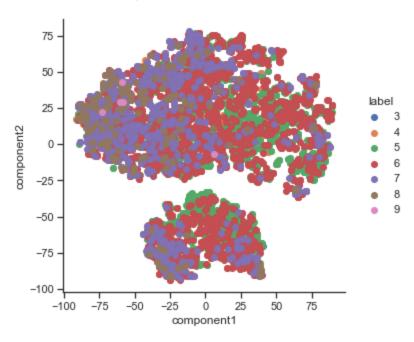
```
LDA Model - Explained Variance of each component
Explained Variance Ratio
0 0.861621
1 0.088287
```



#### T-SNE

```
g = sns.FacetGrid(tsne_df, hue='label', height=5).map(plt.scatter,
    'component1', 'component2')
g.add_legend()
g.fig.suptitle("First Two Components from T-SNE of wine dataset")
g.tight_layout()
```

#### First Two Components from T-SNE of wine dataset

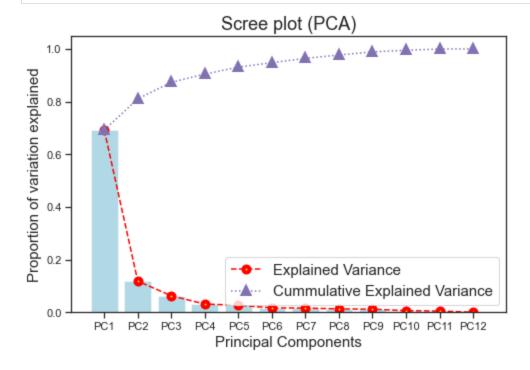


#### Comments

- PCA:Seperation on X-axis(PC1) is more significant and two clusters are formed, spread among each cluster along X-axis is less than Y-axis (color is categorical variable and could be the reason why two different clusters are formed), we have not removed outliers and hence does not perform well.
- LDA: Same class labels are clustered together and is able to better classify between different classes (number of datapoints in various classes are highly different)
- t-SNE: used default settings, non-linear function, clusters similar classes as distance between them are less in both the clusters

#### Scree Plot (PCA)

```
,linewidth=1.5,label='Cummulative Explained Variance')
plt.ylabel('Proportion of variation explained',fontsize = 15)
plt.xlabel('Principal Components',fontsize = 15)
plt.title('Scree plot (PCA)',fontsize = 18)
plt.legend(loc='best',fontsize = 15)
#plt.axis("off")
plt.tight_layout()
plt.show()
```



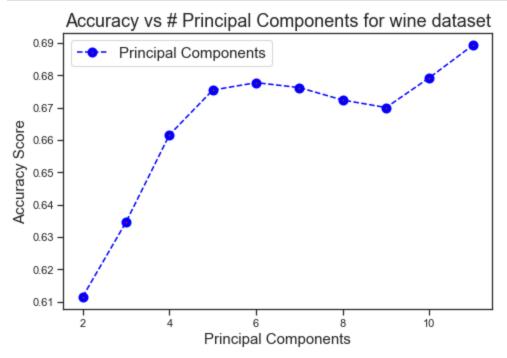
## **Reduced Dimensionality**

#### PCA

```
In [11]:
         knn = KNeighborsClassifier(n neighbors=23 , metric = 'manhattan' , weights =
         "distance")
         D = data.shape[1]
         #print(D)
         acc list =[]
         for i in range (2, D):
             pca = PCA(n components=i)
             X train r = pca.fit(X train).transform(X train)
             X test r = pca.transform(X test)
             knn.fit(X train r, Y train)
             y pred = knn.predict(X test r)
             acc list.append(metrics.accuracy score(Y test, y pred))
         fig = plt.figure(figsize=(7,5))
         plt.plot(range(2,D),acc list, color = 'blue', marker ='o', mew =4, mfc='red',ls
         ='--' ,linewidth=1.5,label='Principal Components')
```

```
plt.ylabel('Accuracy Score', fontsize = 15)
plt.xlabel('Principal Components', fontsize = 15)
plt.title('Accuracy vs # Principal Components for wine dataset', fontsize =
18)
plt.legend(loc='best', fontsize = 15)
plt.tight_layout()
plt.show()

print("The maximum accuracy {} happens to be when number of Principal
Components is : {}".format(max(acc_list), acc_list.index(max(acc_list))+2))
```

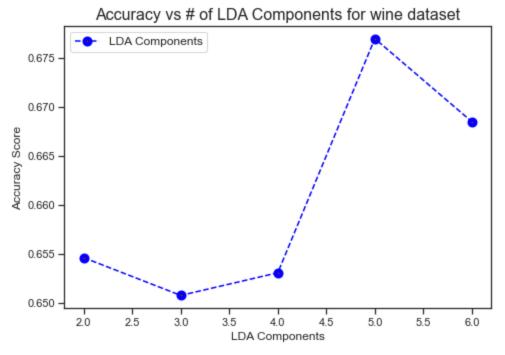


The maximum accuracy 0.6892307692307692 happens to be when number of Principal Components is: 11

#### LDA

```
fig = plt.figure(figsize=(7,5))
plt.plot(range(2,D),acc_list, color = 'blue',marker ='o',mew =4,mfc='red',ls
='--' ,linewidth=1.5,label='LDA Components')
plt.ylabel('Accuracy Score',fontsize = 12)
plt.xlabel('LDA Components',fontsize = 12)
plt.title('Accuracy vs # of LDA Components for wine dataset',fontsize = 16)
plt.legend(loc='best',fontsize = 12)
plt.tight_layout()
plt.show()

print("The maximum accuracy {} happens to be when number of LDA Components is
: {}".format(max(acc_list),acc_list.index(max(acc_list))+2))
```



The maximum accuracy 0.676923076923077 happens to be when number of LDA Components is: 5

#### Comments

- Have used 5 principal components in subsequent parts as there is not a significant difference when included more components. Also, 11 components is the original dimension of data and hence would be better representation but there is no significant loss of accuracy by including 5 components.
- LDA gives better accuracy with 5 components.
- Accuracy with reduced dimensions is comparable with accuracy from Assignment-1 (0.68). PCA performs slightly better with 0.69 accuracy. The major observation is that with reduced dimensions accuracy is comparable with the original datset(with all features) hence PCA an dLDA are able to capture most information in lesser components.

### Versions of dataset (raw/pca/lda)

```
pca = PCA(n components=5)
X r = pca.fit(x).transform(x)
pca data = np.vstack((X r.T, target)).T
wine pca = pd.DataFrame(data = pca data, columns =
['component1','component2','component3', 'component4','component5','label'] )
wine pca temp = wine pca.copy()
wine pca temp 2 = wine pca temp.drop(columns = ['label'])
wine pca = (wine pca temp 2 - wine pca temp 2.min(axis=0)) / (
wine pca temp 2.max(axis=0) - wine pca temp 2.min(axis=0))
wine pca['label']=wine pca temp['label']
wine pca['label'] = wine pca['label'].astype('int')
wine pca['label'] = wine pca['label'].astype('category')
#print(wine pca.head())
lda = LinearDiscriminantAnalysis(n components=5)
X r2 = lda.fit(x, y).transform(x)
lda data = np.vstack((X r2.T, target)).T
wine lda = pd.DataFrame(data = lda data, columns = ['component1',
'component2','component3','component4','component5','label'] )
wine lda temp = wine lda.copy()
wine lda temp 2 = wine lda temp.drop(columns = ['label'])
wine 1da = (wine 1da temp 2 - wine 1da temp 2.min(axis=0)) / (
wine lda temp 2.max(axis=0) - wine lda temp 2.min(axis=0))
wine lda['label']=wine lda temp['label']
wine lda['label'] = wine lda['label'].astype('int')
wine lda['label'] = wine lda['label'].astype('category')
#print(wine lda.head())
```

```
In [14]: wine_p = wine_pca.copy()
  target_pca = wine_p['label']
  data_pca = wine_p.drop(columns = 'label')
  X_train_pca, X_test_pca, Y_train_pca, Y_test_pca = train_test_split(data_pca,
  target_pca, test_size=0.2 , random_state=27)

wine_l = wine_lda.copy()
  target_lda = wine_l['label']
  data_lda = wine_l.drop(columns = 'label')
  X_train_lda, X_test_lda, Y_train_lda, Y_test_lda = train_test_split(data_lda,
  target_lda, test_size=0.2 , random_state=27)
```

## **Naive Bayes Classifier**

5-Fold CV for Naive Bayes & KNN classifier

```
In [15]:
       kfold = KFold(n splits=5)
        clrs = []
        clrs.append(KNeighborsClassifier(n neighbors=23 , metric = 'manhattan' ,
        weights = "distance"))
        clrs.append(GaussianNB())
        clrs.append(MultinomialNB())
        clrs.append(ComplementNB())
        cv results = []
        cv results pca = []
        cv results lda = []
        for clr in clrs :
            cv results.append(cross val score(clr, X train, Y train, scoring =
        'accuracy', cv = kfold, n jobs=-1))
            cv results pca.append(cross val score(clr, X train pca, Y train pca,
        scoring = 'accuracy', cv = kfold, n jobs=-1))
            cv results lda.append(cross val score(clr, X train lda, Y train lda,
        scoring = 'accuracy', cv = kfold, n jobs=-1))
        cv means = []
        cv means pca = []
        cv means lda = []
        for cv result in cv results:
            cv means.append(cv result.mean())
        for cv result in cv results pca:
            cv means pca.append(cv result.mean())
        for cv result in cv results lda:
            cv means lda.append(cv result.mean())
        cv df = pd.DataFrame({"Mean Accuracy":cv means,"Algo":
        ['KNeighborsClassifier', 'GaussianNB', 'MultinomialNB', 'ComplementNB']})
        cv df pca = pd.DataFrame({"Mean Accuracy":cv means pca, "Algo":
        ['KNeighborsClassifier','GaussianNB','MultinomialNB','ComplementNB']})
        cv df lda = pd.DataFrame({"Mean Accuracy":cv means lda,"Algo":
        ['KNeighborsClassifier','GaussianNB','MultinomialNB','ComplementNB']})
         #print(cv df)
         #print(cv df pca)
         #print(cv df lda)
```

```
In [16]:
        data frames = [cv df, cv df pca, cv df lda]
        df merged = reduce(lambda left, right: pd.merge(left, right, on=
         ['Algo'], how='outer'), data frames)
        df merged.rename(columns={"Mean Accuracy x": "wine raw", "Mean Accuracy y":
        "wine pca", "Mean Accuracy": "wine lda"}, inplace=True)
        df merged.set index('Algo', inplace=True)
        print(df merged)
                            wine raw wine pca wine lda
       Algo
       KNeighborsClassifier 0.652869 0.640555 0.646522
                            0.359820 0.509904 0.537423
       GaussianNB
       MultinomialNB
                           0.472769 0.436018 0.436018
       ComplementNB
                           0.396376 0.367124 0.438711
In [17]:
        kfold = KFold(n splits=5)
        clrs = []
        clrs.append(KNeighborsClassifier(n neighbors=23, metric = 'manhattan',
        weights = "distance"))
        clrs.append(GaussianNB())
        clrs.append(MultinomialNB())
        clrs.append(ComplementNB())
        cv results = []
        cv results pca = []
        cv results lda = []
         #scoring = {'accuracy' : make scorer(accuracy score),
                     'precision' : make scorer(precision score),
                     'recall' : make scorer(recall score),
                    'f1 score' : make scorer(f1 score)}
        for clr in clrs :
            cv results.append(cross val score(clr, data, target , scoring =
         'accuracy', cv = kfold, n jobs=-1))
             cv results pca.append(cross val score(clr, data pca, target pca, scoring
        = 'accuracy', cv = kfold, n jobs=-1))
             cv results lda.append(cross val score(clr, data pca, target pca, scoring
        = 'accuracy', cv = kfold, n jobs=-1))
        cv means = []
        cv means pca = []
        cv means lda = []
```

```
#print(cv results)
for cv result in cv results:
    cv means.append(cv result.mean())
for cv result in cv results pca:
    cv means pca.append(cv result.mean())
for cv result in cv results lda:
    cv means lda.append(cv result.mean())
cv df = pd.DataFrame({"Mean Accuracy":cv means,"Algo":
['KNeighborsClassifier','GaussianNB','MultinomialNB','ComplementNB']})
cv df pca = pd.DataFrame({"Mean Accuracy":cv means pca,"Algo":
['KNeighborsClassifier','GaussianNB','MultinomialNB','ComplementNB']})
cv df lda = pd.DataFrame({"Mean Accuracy":cv means lda,"Algo":
['KNeighborsClassifier','GaussianNB','MultinomialNB','ComplementNB']})
#print(cv df)
#print(cv df pca)
#print(cv df lda)
```

```
In [18]: data_frames = [cv_df, cv_df_pca, cv_df_lda]
    df_merged = reduce(lambda left,right: pd.merge(left,right,on=
        ['Algo'],how='outer'), data_frames)
    df_merged.rename(columns={"Mean_Accuracy_x": "wine_raw", "Mean_Accuracy_y":
        "wine_pca","Mean_Accuracy":"wine_lda"},inplace=True)
    df_merged.set_index('Algo',inplace=True)
    print(df_merged)
```

```
wine_rawwine_pcawine_ldaAlgo0.5042360.4893100.489310GaussianNB0.3095050.4671390.467139MultinomialNB0.4303560.4365150.436515ComplementNB0.3963360.3854060.385406
```

#### **Comments**

• Complement Naive Bayes does not outperform as expected for the unbalance wine dataset. While training using Complement NB it trains each class with all data, but the sample from that class and because it is a multiclass classification with higer Prior, the likelihood impact is reduced and overall accuracy is not good.

## **Decision Trees Classifier**

## 5-Fold CV and hyperparameter tuning

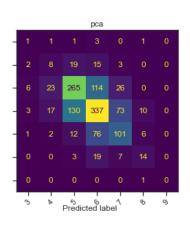
```
dtc = DecisionTreeClassifier()
dtc.fit(X train, Y train)
Y pred = dtc.predict(X test)
#cm=confusion matrix(Y test, Y pred)
#print(confusion matrix(Y test, Y pred))
#print(classification report(Y test, Y pred))
print ("Accuracy Score for Decision Tree classifier without hyper parameter
tuning is: ",accuracy score(Y test, Y pred))
\#disp =
ConfusionMatrixDisplay(confusion matrix=cm, display labels=dtc.classes)
#disp.plot()
grid params = {
    'criterion' : ['gini', 'entropy'],
    'max depth' : range(2,25,1),
    'min samples split' : range(2, 10, 1),
    'min samples leaf' : range(2, 10, 1)
grid search = GridSearchCV(dtc, grid params, cv = 5, n jobs = -1, verbose =
1)
grid search.fit(X train, Y train)
dtc = grid search.best estimator
Y pred = dtc.predict(X test)
print("Accuracy Score is :",accuracy_score(Y_test, Y_pred))
# best parameters and best score
print("Best Parameters are : ", grid search.best params )
print("Best Score is : ",grid search.best score )
#dt raw acc = accuracy score(Y test, Y pred)
dt raw set=grid search.best params
grid search.fit(X train pca, Y train pca)
dtc = grid search.best estimator
Y pred pca = dtc.predict(X test pca)
print("Accuracy Score using PCA is :",accuracy score(Y test pca, Y pred pca))
# best parameters and best score
print("Best Parameters using PCA are : ",grid search.best params )
print("Best Score using PCA is : ",grid search.best score )
dt pca set=grid search.best params
grid_search.fit(X_train_lda, Y train lda)
dtc = grid search.best estimator
Y pred lda = dtc.predict(X test lda)
print ("Accuracy Score using LDA is : ", accuracy score (Y test lda, Y pred lda))
```

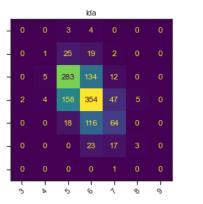
```
# best parameters and best score
         print("Best Parameters using LDA are : ", grid search.best params )
         print("Best Score using LDA is : ", grid search.best score )
         dt lda set=grid search.best params
        Accuracy Score for Decision Tree classifier without hyper parameter tuning is: 0.61153846
        15384616
        Fitting 5 folds for each of 2944 candidates, totalling 14720 fits
        Accuracy Score is : 0.5792307692307692
        Best Parameters are : {'criterion': 'entropy', 'max depth': 18, 'min samples leaf': 2, 'm
        in samples split': 6}
        Best Score is : 0.5639766417413193
        Fitting 5 folds for each of 2944 candidates, totalling 14720 fits
        Accuracy Score using PCA is : 0.5584615384615385
        Best Parameters using PCA are : {'criterion': 'entropy', 'max depth': 18, 'min samples le
        af': 2, 'min samples split': 2}
        Best Score using PCA is: 0.5474289257422077
        Fitting 5 folds for each of 2944 candidates, totalling 14720 fits
        Accuracy Score using LDA is : 0.5423076923076923
        Best Parameters using LDA are : {'criterion': 'gini', 'max depth': 8, 'min samples leaf':
        2, 'min samples split': 6}
        Best Score using LDA is : 0.5505084400681128
In [20]:
        dt raw acc = accuracy score(Y test, Y pred)
         dt pca acc = accuracy_score(Y_test_pca, Y_pred_pca)
         dt lda acc = accuracy score (Y test lda, Y pred lda)
         #print(dt raw acc,dt pca acc,dt lda acc)
In [21]:
         cm=confusion matrix(Y test, Y pred)
         cm pca=confusion matrix(Y test pca, Y pred pca)
         cm lda=confusion matrix(Y_test_lda, Y_pred_lda)
         cm dict = {
             "raw": cm,
             "pca": cm pca,
             "lda": cm lda
         f, axes = plt.subplots(1, 3, figsize=(20, 5), sharey='row')
         for i, (c,cm) in enumerate(cm dict.items()):
             #print(i,cm)
             disp = ConfusionMatrixDisplay(cm, display labels=dtc.classes )
             disp.plot(ax=axes[i], xticks rotation=45)
             disp.ax .set title(c)
             disp.im .colorbar.remove()
             disp.ax .set xlabel('')
             if i!=0:
                 disp.ax .set ylabel('')
```

```
f.text(0.4, 0.1, 'Predicted label', ha='left')
plt.subplots_adjust(wspace=0.40, hspace=0.1)

f.suptitle('Confusion matrix', fontsize=12)
f.colorbar(disp.im_, ax=axes)
plt.show()
```

Confusion matrix



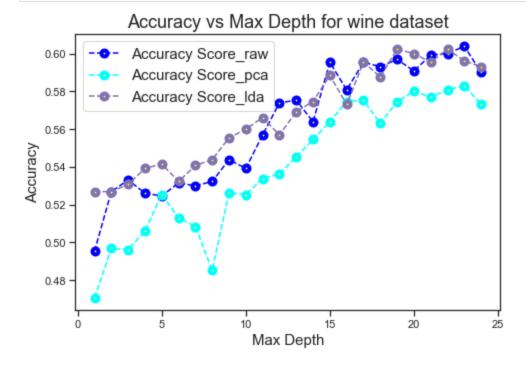


# - 350 - 300 - 250 - 200 - 150 - 100 - 50

## Accuracy vs Max\_depth

```
In [22]:
        def plot acc vs depth(dataset):
            max depth range = list(range(1, 25))
             acc list = []
             acc list pca = []
            acc list lda = []
             for depth in max depth range:
                 dtc w = DecisionTreeClassifier(max depth = depth)
                 if dataset[0] == 'raw':
                     dtc w.fit(X train, Y train)
                     score = dtc w.score(X test, Y test)
                     acc list.append(score)
                 if dataset[1] == 'pca':
                     dtc w.fit(X train pca, Y train pca)
                     score = dtc w.score(X test pca, Y test pca)
                     acc list pca.append(score)
                 if dataset[2] == 'lda':
                     dtc w.fit(X train lda, Y train lda)
                     score =dtc w.score(X test lda, Y test lda)
                     acc list lda.append(score)
             return (acc list, acc list pca, acc list lda)
        acc list,acc list pca,acc list_lda = plot_acc_vs_depth(['raw','pca','lda'])
        fig = plt.figure(figsize=(7,5))
        plt.plot(range(1,25),acc list, color = 'blue',marker ='o',mew
```

```
=4, mfc='yellow', ls ='--', linewidth=1.5, label='Accuracy Score_raw')
plt.plot(range(1,25),acc_list_pca, color = 'cyan', marker ='o', mew
=4, mfc='yellow', ls ='--', linewidth=1.5, label='Accuracy Score_pca')
plt.plot(range(1,25),acc_list_lda, color = 'm', marker ='o', mew
=4, mfc='yellow', ls ='--', linewidth=1.5, label='Accuracy Score_lda')
plt.ylabel('Accuracy', fontsize = 15)
plt.xlabel('Max Depth', fontsize = 15)
plt.title('Accuracy vs Max Depth for wine dataset', fontsize = 18)
plt.legend(loc='best', fontsize = 15)
plt.tight_layout()
plt.show()
```



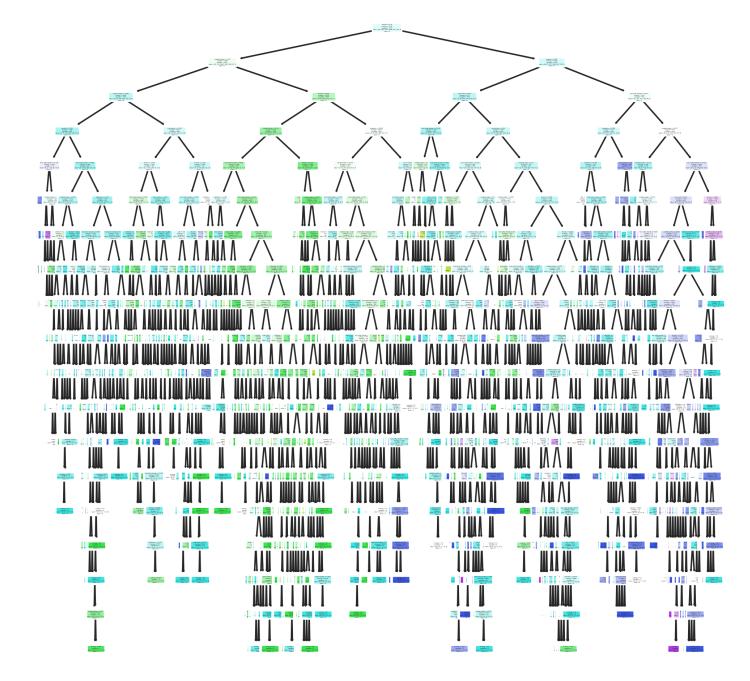
#### Splitting rules used for the trees

## Text split rules

In [8]: from sklearn.tree import export\_text

```
dtc_g = DecisionTreeClassifier(criterion= 'entropy', max_depth=18,
min_samples_leaf= 2, min_samples_split=5)
dt_rules = dtc_g.fit(X_train,Y_train)
fn_t=X_train.columns
#print(type(fn_t.tolist()))
tree_rules = export_text(dt_rules, feature_names = fn_t.tolist())
#print(tree_rules)
```

#### Splitting rules used for the trees but fitted on page



#### **Comments**

• Alcohol and volatile accidity are important splitting parametrs. Optimal Max\_depth is larger than the original features, clearly indicating that there are no clear patterns while splitting. Categorical and quantitave, both data are used for splitting rules.

## **Random Forest Classifier**

## 5-Fold CV and hyperpararmeter tuning

```
print ("Accuracy Score for Forest classifier without hyper parameter tuning
is: ",accuracy score(Y test, Y pred))
#disp =
ConfusionMatrixDisplay(confusion matrix=cm, display labels=dtc.classes )
#disp.plot()
grid params = {
    'n estimators': [2,5,200, 500],
    'criterion' : ['entropy'],
    'max depth' : range(2,25,1),
    #'max features': range(3,10,1),
    #'min samples split' : range(2, 10, 1),
    #'min samples leaf' : range(2, 10, 1)
grid search = GridSearchCV(rfc, grid params, cv = 5, n jobs = -1, verbose =
True)
#print(grid search.get params().keys())
grid search.fit(X train, Y train)
rfc = grid search.best estimator
Y pred = rfc.predict(X test)
print("Accuracy Score is :",accuracy score(Y test, Y pred))
rf raw acc=accuracy score(Y test, Y pred)
# best parameters and best score
print("Best Parameters are : ", grid search.best params )
rf raw set=grid search.best params
#print("Best Score is : ", grid search.best score )
ac df=pd.DataFrame(grid search.cv results ['params'])
#Creating a data frame with hyperparameters and accuracy
ac df["accuracy"]=grid_search.cv_results_['mean_test_score']
grid search.fit(X train pca, Y train pca)
rfc pca = grid search.best estimator
Y pred pca = rfc pca.predict(X test pca)
print("Accuracy Score using PCA is :",accuracy score(Y test pca, Y pred pca))
rf pca acc=accuracy score(Y test pca, Y pred pca)
# best parameters and best score
print("Best Parameters using PCA are : ", grid search.best params )
rf pca set=grid search.best params
#print("Best Score using PCA is : ",grid search.best score )
ac df pca=pd.DataFrame(grid search.cv results ['params'])
#Creating a data frame with hyperparameters and accuracy
ac df pca["accuracy"]=grid search.cv results ['mean test score']
```

```
grid_search.fit(X_train_lda, Y_train_lda)
rfc_lda = grid_search.best_estimator_
Y_pred_lda = rfc_lda.predict(X_test_lda)
print("Accuracy Score using LDA is :",accuracy_score(Y_test_lda, Y_pred_lda))
rf_lda_acc=accuracy_score(Y_test_lda, Y_pred_lda)
# best parameters and best score
print("Best Parameters using LDA are : ",grid_search.best_params_)
#print("Best Score using LDA is : ",grid_search.best_score_)
rf_lda_set=grid_search.best_params_
ac_df_lda=pd.DataFrame(grid_search.cv_results_['params'])
#Creating a data frame with hyperparameters and accuracy
ac_df_lda["accuracy"]=grid_search.cv_results_['mean_test_score']
```

```
Accuracy Score for Forest classifier without hyper parameter tuning is: 0.6915384615384615

Fitting 5 folds for each of 92 candidates, totalling 460 fits

Accuracy Score is: 0.6953846153846154

Best Parameters are: {'criterion': 'entropy', 'max_depth': 18, 'n_estimators': 500}

Fitting 5 folds for each of 92 candidates, totalling 460 fits

Accuracy Score using PCA is: 0.6607692307692308

Best Parameters using PCA are: {'criterion': 'entropy', 'max_depth': 16, 'n_estimators': 200}

Fitting 5 folds for each of 92 candidates, totalling 460 fits

Accuracy Score using LDA is: 0.6784615384615384

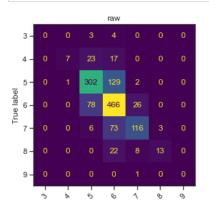
Best Parameters using LDA are: {'criterion': 'entropy', 'max_depth': 17, 'n_estimators': 500}
```

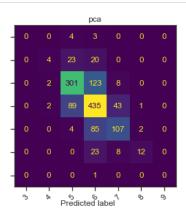
In [26]:

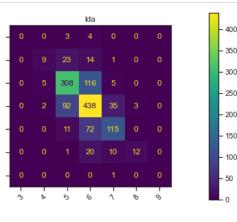
```
cm=confusion matrix(Y test, Y pred)
cm pca=confusion matrix(Y_test_pca, Y_pred_pca)
cm lda=confusion matrix(Y test lda, Y pred lda)
cm dict = {
    "raw": cm,
    "pca": cm pca,
    "lda": cm lda
}
f, axes = plt.subplots(1, 3, figsize=(20, 5), sharey='row')
for i, (c,cm) in enumerate(cm dict.items()):
    #print(i,cm)
    disp = ConfusionMatrixDisplay(cm, display labels=dtc.classes )
    disp.plot(ax=axes[i], xticks rotation=45)
    disp.ax .set title(c)
    disp.im .colorbar.remove()
    disp.ax .set xlabel('')
     if i!=0:
         disp.ax .set ylabel('')
```

```
f.text(0.4, 0.1, 'Predicted label', ha='left')
plt.subplots_adjust(wspace=0.40, hspace=0.1)

f.colorbar(disp.im_, ax=axes)
plt.show()
```







## Heatmap of max\_depth vs n\_estimators

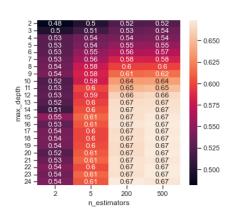
```
In [27]:
```

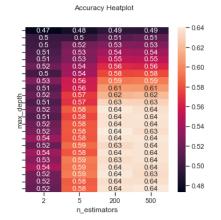
```
#Pivoting the dataframe for plotting heat map
ac_df=ac_df.pivot(index='max_depth',columns='n_estimators',values='accuracy')
ac_df_pca=ac_df_pca.pivot(index='max_depth',columns='n_estimators',values='accuracy')
ac_df_lda=ac_df_lda.pivot(index='max_depth',columns='n_estimators',values='accuracy')

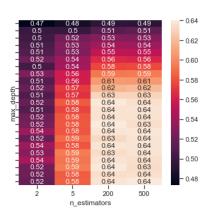
#Plotting the graph
fig, ax =plt.subplots(1,3,figsize=(20, 5), sharey='row')

sns.heatmap(ac_df,annot=True, ax=ax[0])
sns.heatmap(ac_df_pca,annot=True, ax=ax[1])
sns.heatmap(ac_df_pca,annot=True, ax=ax[2])

plt.subplots_adjust(wspace=0.40, hspace=0.1)
fig.suptitle('Accuracy Heatplot', fontsize=12)
plt.show()
```







## **Gradient Boosting Classifier**

#### 5-Fold CV and hyperparameter tuning

```
In [28]:
        gbc = GradientBoostingClassifier()
        gbc.fit(X train, Y train)
        Y pred = gbc.predict(X test)
        cm=confusion matrix(Y test, Y pred)
        #print(confusion matrix(Y test, Y pred))
        #print(classification report(Y test, Y pred))
        print ("Accuracy Score for Gradient Boosting classifier without hyper
        parameter tuning is: ",accuracy score(Y test, Y pred))
        ConfusionMatrixDisplay(confusion matrix=cm, display labels=dtc.classes)
        #disp.plot()
        grid params = {
             'n estimators': [2,120,200,500,800],
             'learning rate' : [0.01, 0.1,1],
             #'criterion' : [None, 'mse', 'mae'],
            #'max depth' : range(2,22,2),
            #'max features': range(3,10,1),
            #'min samples split' : range(2, 10, 1),
             #'min samples leaf' : range(2, 10, 1)
        grid search = GridSearchCV(gbc, grid params, cv = 5, n jobs = -1, verbose =
        #print(grid search.get params().keys())
        grid search.fit(X_train, Y_train)
        gbc = grid search.best estimator
        Y pred = gbc.predict(X test)
        print("Accuracy Score is :",accuracy score(Y test, Y pred))
        gbc acc=accuracy score(Y test, Y pred)
        # best parameters and best score
        print("Best Parameters are : ", grid search.best params )
        gbc set=grid search.best params
        #print("Best Score using PCA is : ",grid search.best score )
        gbc ac df=pd.DataFrame(grid search.cv results ['params'])
        #Creating a data frame with hyperparameters and accuracy
        gbc ac df["accuracy"]=grid search.cv results ['mean test score']
```

```
grid search.fit(X train pca, Y train pca)
 gbc = grid search.best estimator
 Y pred pca = gbc.predict(X test pca)
 print("Accuracy Score is :",accuracy score(Y test pca, Y pred pca))
 gbc pca acc=accuracy_score(Y_test_pca, Y_pred_pca)
 # best parameters and best score
 print("Best Parameters are : ", grid search.best params )
 gbc pca set=grid search.best params
 #print("Best Score using PCA is : ",grid search.best score )
 gbc ac df pca=pd.DataFrame(grid search.cv results ['params'])
 #Creating a data frame with hyperparameters and accuracy
 gbc ac df pca["accuracy"]=grid search.cv results ['mean test score']
 grid search.fit(X train lda, Y train lda)
 gbc = grid search.best estimator
Y pred lda = gbc.predict(X test lda)
 print("Accuracy Score is :", accuracy score(Y test lda, Y pred lda))
 gbc lda acc=accuracy score(Y test lda, Y pred lda)
 # best parameters and best score
 print("Best Parameters are : ", grid search.best params )
 gbc lda set=grid search.best params
 #print("Best Score using PCA is : ",grid search.best score )
 gbc ac df lda=pd.DataFrame(grid search.cv results ['params'])
 #Creating a data frame with hyperparameters and accuracy
 gbc ac df lda["accuracy"]=grid search.cv results ['mean test score']
Accuracy Score for Gradient Boosting classifier without hyper parameter tuning is: 0.6007
692307692307
Fitting 5 folds for each of 15 candidates, totalling 75 fits
```

```
692307692307

Fitting 5 folds for each of 15 candidates, totalling 75 fits

Accuracy Score is: 0.6346153846153846

Best Parameters are: {'learning_rate': 0.1, 'n_estimators': 800}

Fitting 5 folds for each of 15 candidates, totalling 75 fits

Accuracy Score is: 0.6192307692307693

Best Parameters are: {'learning_rate': 0.1, 'n_estimators': 800}

Fitting 5 folds for each of 15 candidates, totalling 75 fits

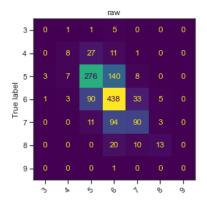
Accuracy Score is: 0.6430769230769231

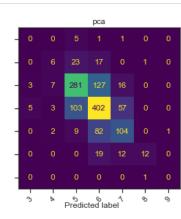
Best Parameters are: {'learning_rate': 0.1, 'n_estimators': 800}
```

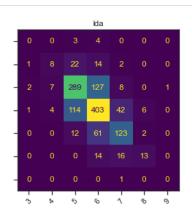
```
f, axes = plt.subplots(1, 3, figsize=(20, 5), sharey='row')
for i,(c,cm) in enumerate(cm_dict.items()):
    #print(i,cm)
    disp = ConfusionMatrixDisplay(cm,display_labels=dtc.classes_)
    disp.plot(ax=axes[i], xticks_rotation=45)
    disp.ax_.set_title(c)
    disp.im_.colorbar.remove()
    disp.ax_.set_xlabel('')
    if i!=0:
        disp.ax_.set_ylabel('')

f.text(0.4, 0.1, 'Predicted label', ha='left')
plt.subplots_adjust(wspace=0.40, hspace=0.1)

f.colorbar(disp.im_, ax=axes)
plt.show()
```







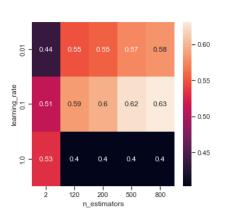
- 300

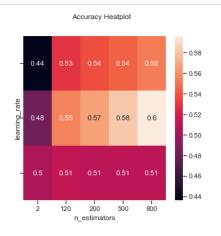
- 200

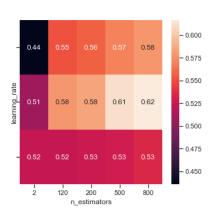
# Heatplot of learning\_rate and n\_estimators

```
In [30]: #Pivoting the dataframe for plotting heat map
  gbc_ac_df=gbc_ac_df.pivot(index='learning_rate',columns='n_estimators',values=
  gbc_ac_df_pca=gbc_ac_df_pca.pivot(index='learning_rate',columns='n_estimators'
  gbc_ac_df_lda=gbc_ac_df_lda.pivot(index='learning_rate',columns='n_estimators'
  #Plotting the graph
  fig, ax =plt.subplots(1,3,figsize=(20, 5), sharey='row')
  sns.heatmap(gbc_ac_df,annot=True, ax=ax[0])
  sns.heatmap(gbc_ac_df_pca,annot=True,ax=ax[1])
  sns.heatmap(gbc_ac_df_lda,annot=True,ax=ax[2])
```

```
plt.subplots_adjust(wspace=0.40, hspace=0.1)
fig.suptitle('Accuracy Heatplot', fontsize=12)
plt.show()
```







### **Final Results**

```
Algo
                       wine_raw wine_pca wine_lda
   KNeighborsClassifier
                        0.504236
                                  0.489310
                                             0.489310
1
          GaussianNB
                        0.309505
                                  0.467139
                                             0.467139
2
        MultinomialNB
                        0.430356
                                  0.436515
                                             0.436515
3
                                             0.385406
       ComplementNB
                        0.396336
                                  0.385406
         Decision Tree
                        0.579231
                                  0.558462
                                             0.542308
4
5
        Random Forest
                        0.695385
                                  0.660769
                                             0.678462
6
     Gradient Boosting
                        0.634615
                                  0.619231 0.643077
```

```
In [ ]: #print(dt_raw_set.keys())
    #print(dt_raw_set.values(),dt_pca_set.values(),dt_lda_set.values())
    dt_s =[dt_raw_set,dt_pca_set,dt_lda_set]
    df_dt_settings = pd.DataFrame(dt_s)
    df_dt_settings['Algo'] = 'Decision Tree'
```

```
df_dt_settings['Data'] = ['raw', 'pca', 'lda']
display(df_dt_settings)
```

```
criterionmax_depthn_estimatorsAlgoData0entropy18500Random Forestraw1entropy16200Random Forestpca2entropy17500Random ForestIda
```

```
In [66]: gbc_s =[gbc_set,gbc_pca_set,gbc_lda_set]
    df_gbc_settings = pd.DataFrame(gbc_s)
    df_gbc_settings['Algo'] = 'Gradient Boosting'
    df_gbc_settings['Data'] = ['raw','pca','lda']
    display(df_gbc_settings)
```

```
learning_raten_estimatorsAlgoData00.1800Gradient Boostingraw10.1800Gradient Boostingpca20.1800Gradient BoostingIda
```

```
In []: knn_setting = ({'n_neighbors':23 , 'metric': 'manhattan' , 'weights':
    "distance"})
    gnb_settings = ("default")
    mnb_settings = ("default")
    dt_settings = ("default")
    dt_settings =
        (('criterion':'Entropy', 'max_depth':18, 'min_samples_leaf':2, 'min_samples_split')

    rf_settings = ({'criterion':'Entropy', 'max_depth':18, 'n_estimators':500})
    gb_settings = ({'learning_rate':0.1, 'n_estimators':800})

    settings_all =
        [knn_setting,gnb_settings,mnb_settings,cnb_settings,dt_settings,rf_settings_all
        df_final['settings'] =settings_all
```

	Algo	settings	wine_raw	wine_pca	wine_lda
0	KNeighborsClassifier	('n_neighbors': 23, 'metric': 'manhattan', 'we	0.504236	0.489310	0.489310
1	GaussianNB	default'	0.309505	0.467139	0.467139
2	MultinomialNB	default'	0.430356	0.436515	0.436515
3	ComplementNB	default'	0.396336	0.385406	0.385406
4	Decision Tree	{'criterion': 'Entropy','max_depth': 18,'min_s	0.579231	0.558462	0.542308
5	Random Forest	{'criterion': 'Entropy', 'max_depth': 18, 'n_e	0.695385	0.660769	0.678462
6	Gradient Boosting	{'learning_rate': 0.1, 'n_estimators': 800}	0.634615	0.619231	0.643077

#### Comments

- Random Forest with following settings *criterion*: entropy, *max-depth*: 18, *n-estimators*:500 on raw data(MinMax Scaling) as well as for PCA and LDA dataset performed the best task at classification.
- For Gradient Boosting and Gaussian Naive bayes, dimensionality reduction did help while for other algorithms the accuracy results were almost comparable to original dataset

## **Abalone Dataset**

## **Preprocessing**

#### **Loading Data**

#### Normalization/Scaling

```
In [29]: abalone1=abalone.copy()
   abalone_z = abalone1.drop(columns = ['Sex', 'Rings'])
   abalone_znormalized = abalone_z.apply(stats.zscore)
   abalone_znormalized['Sex'] = abalone['Sex']
   abalone_znormalized['Rings'] = abalone['Rings']

target_z = abalone_znormalized['Rings']
   data_z = abalone_znormalized.drop(columns = 'Rings')
   X_train_z, X_test_z, Y_train_z, Y_test_z = train_test_split(data_z, target_z, test_size=0.2 , random_state=27)
```

#### Best formulated KNN from Assignment-1

```
In [31]: knn = KNeighborsClassifier(n_neighbors=66, metric='minkowski',
    weights='uniform', p=2)
    knn.fit(X_train, Y_train)
    Y_pred = knn.predict(X_test)

knn_train_acc = knn.score(X_train, Y_train)
    print("Training Score: ", knn_train_acc)
    knn_test_acc = knn.score(X_test, Y_test)
    print("Testing Score: ", knn_test_acc)
```

Training Score: 0.2975157138581263 Testing Score: 0.2631578947368421

## Representation Learning

#### Independent and dependent variable

```
In [32]: #Scaled 'data' and 'target'
    x=data
    y=target
    n_components=2
    target_names = np.sort(y.unique())
```

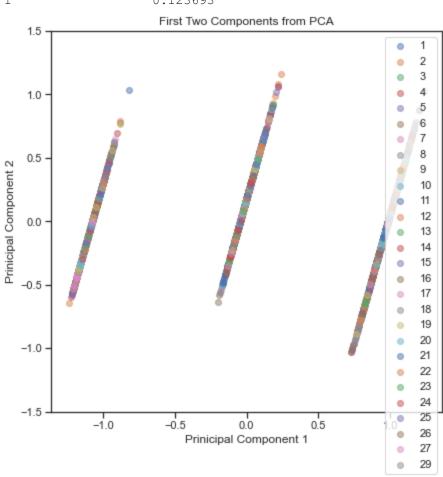
#### **PCA**

```
In [33]:
        print("PCA Model - Explained Variance of each component")
        pca = PCA(n components=n components)
        X r = pca.fit(x).transform(x)
        exp var ratio df = pd.DataFrame(pca.explained variance ratio, columns=
        ['Explained Variance Ratio'])
        print(exp var ratio df)
        plt.figure(figsize=(7,7))
        colors = plt.cm.get cmap("twilight")
        for i, target name in enumerate(target names):
            plt.scatter(
                 X r[y == i, 0], X r[y == i, 1], alpha=0.5, lw=1, label=target name,
        cmap="twilight"
        plt.ylim(-1.5, 1.5)
        plt.legend(loc="best", shadow=False, scatterpoints=1)
        plt.xlabel("Prinicipal Component 1")
        plt.ylabel("Prinicipal Component 2")
```

```
plt.title("First Two Components from PCA")
plt.set_cmap("twilight")

plt.show()
```

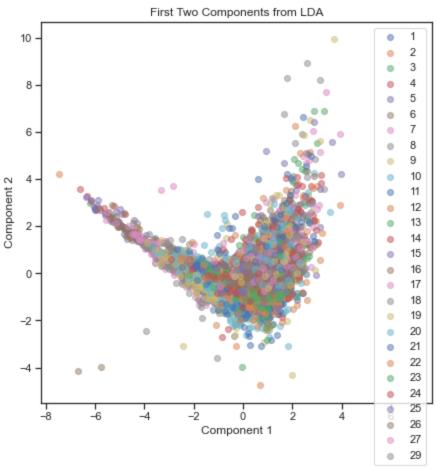
```
PCA Model - Explained Variance of each component
Explained Variance Ratio
0 0.865084
1 0.123693
```



#### LDA

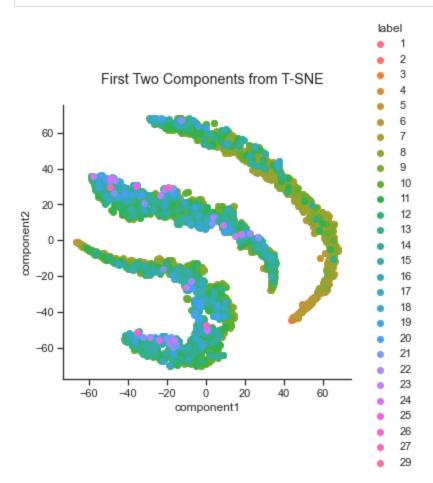
```
#plt.xlim(-4,4)
plt.legend(loc="best", shadow=False, scatterpoints=1)
plt.xlabel("Component 1")
plt.ylabel("Component 2")
plt.title("First Two Components from LDA")
plt.set_cmap("twilight")
```

```
LDA Model - Explained Variance of each component
Explained Variance Ratio
0 0.719139
1 0.216991
```



#### T-SNE

```
g.add_legend()
g.fig.suptitle("First Two Components from T-SNE ")
g.tight_layout()
```

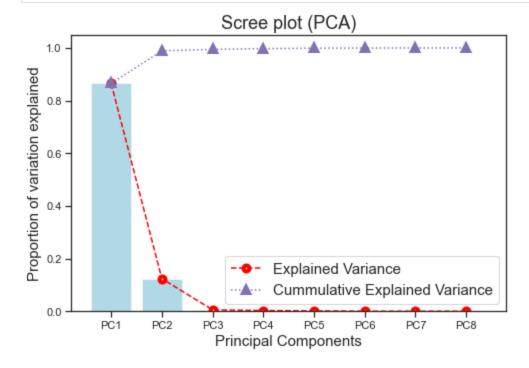


#### Comments

- PCA: Three distant lines due to presence of categorical column(Sex) in the data. We have one principal component that explains nearly all (85%) of the variance the reason is, the principal component is a mixture of all variables in nearly equal proportions. That is, the principal component is the one that combines every single variable.
- LDA: Inter Class spread is increased and intra-class spread is decreased, clusterring similar class points together. Outliers evident in the plot as distance between them is large.
- t-SNE: Keeps changing with the run, three clusters are formed, better way to visualize but doe not help in our case for multiclass classification

### Scree Plot (PCA)

```
,linewidth=1.5,label='Explained Variance')
plt.plot(range(len(explained_variance_pca)), explained_variance_pca.cumsum(),
color = 'm', marker ='^', mew =4, mfc='red', ls =':'
,linewidth=1.5,label='Cummulative Explained Variance')
plt.ylabel('Proportion of variation explained', fontsize = 15)
plt.xlabel('Principal Components', fontsize = 15)
plt.title('Scree plot (PCA)', fontsize = 18)
plt.legend(loc='best', fontsize = 15)
#plt.axis("off")
plt.tight_layout()
plt.show()
```



## **Reduced Dimensionality**

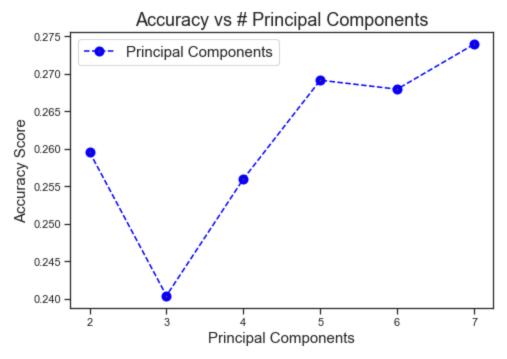
#### PCA

```
In [37]: knn = KNeighborsClassifier(n_neighbors=66, metric='minkowski',
    weights='uniform', p=2)
    D = data.shape[1]
    #print(D)
    acc_list =[]

for i in range(2,D):
    pca = PCA(n_components=i)
    X_train_r = pca.fit(X_train).transform(X_train)
    X_test_r = pca.transform(X_test)
    knn.fit(X_train_r, Y_train)
    y_pred = knn.predict(X_test_r)
    acc_list.append(metrics.accuracy_score(Y_test,y_pred))
```

```
fig = plt.figure(figsize=(7,5))
plt.plot(range(2,D),acc_list, color = 'blue',marker ='o',mew =4,mfc='red',ls
='--' ,linewidth=1.5,label='Principal Components')
plt.ylabel('Accuracy Score',fontsize = 15)
plt.xlabel('Principal Components',fontsize = 15)
plt.title('Accuracy vs # Principal Components',fontsize = 18)
plt.legend(loc='best',fontsize = 15)
plt.tight_layout()
plt.show()

print("The maximum accuracy {} happens to be when number of Principal
Components is : {}".format(max(acc_list),acc_list.index(max(acc_list))+2))
```



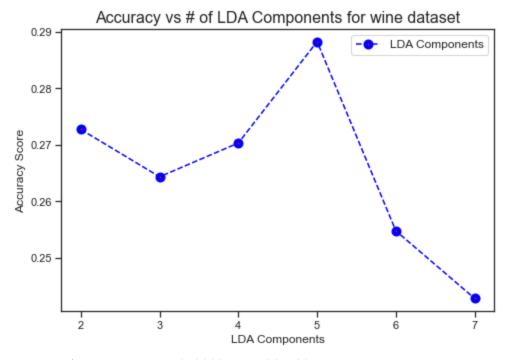
The maximum accuracy 0.27392344497607657 happens to be when number of Principal Components is: 7

#### LDA

```
In [38]: knn = KNeighborsClassifier(n_neighbors=66, metric='minkowski',
    weights='uniform', p=2)
#LDA is constrained by the number of components it can use , given by :
    min(n_classes-1, len(features))
D = min(data.shape[1],len(target_names))
acc_list =[]

for i in range(2,D):
    lda = LinearDiscriminantAnalysis(n_components=i)
    X_train_r = lda.fit(X_train,Y_train).transform(X_train)
    X_test_r = lda.transform(X_test)
    knn.fit(X_train_r, Y_train)
    y_pred = knn.predict(X_test_r)
```

```
acc list.append(metrics.accuracy score(Y test, y pred))
fig = plt.figure(figsize=(7,5))
plt.plot(range(2,D),acc list, color = 'blue', marker ='o', mew =4, mfc='red',ls
='--' ,linewidth=1.5,label='LDA Components')
plt.ylabel('Accuracy Score', fontsize = 12)
plt.xlabel('LDA Components', fontsize = 12)
plt.title('Accuracy vs # of LDA Components for wine dataset', fontsize = 16)
plt.legend(loc='best', fontsize = 12)
plt.tight layout()
plt.show()
print ("The maximum accuracy {} happens to be when number of LDA Components is
  {}".format(max(acc list),acc list.index(max(acc list))+2))
```



The maximum accuracy 0.28827751196172247 happens to be when number of LDA Components is:

#### Comments

- Two Principal Components are able to explain 98% of the variance in the dataset. We get maximum accuracy with 7 principal components which is same as orginal dimension of dataset, but we use 5 components in subsequent sections as there is no significant difference in accuracy with 5 and 7 components.
- With 5 components in LDA, we get maximum accuracy (more than PCA). LDA does a better job at classification task.
- Dimensionality Reduction improves the accuracy for both PCA and LDA compared to best formulation from Assignment-1

#### Versions of dataset (raw/pca/lda)

```
abalone normalized raw = abalone znormalized.copy()
abalone minmax normalized = abalone minmax.copy()
pca = PCA(n components=5)
X r = pca.fit(x).transform(x)
pca data = np.vstack((X r.T, target)).T
abalone pca = pd.DataFrame(data = pca data, columns =
['component1','component2','component3', 'component4','component5','label'] )
abalone temp pca = abalone pca.copy()
abalone_temp_pca_2 = abalone_temp_pca.drop(columns = ['label'])
abalone pca = (abalone temp pca 2 - abalone temp pca 2.min()) / (
abalone temp pca 2.max() - abalone temp pca 2.min())
abalone pca['label'] = abalone temp pca['label']
#abalone pca = (abalone pca - abalone pca.min()) / ( abalone pca.max() -
abalone pca.min())
abalone pca['label'] = abalone pca['label'].astype('int')
abalone pca['label'] = abalone pca['label'].astype('category')
#print(abalone pca.head())
lda = LinearDiscriminantAnalysis(n components=5)
X r2 = lda.fit(x, y).transform(x)
lda data = np.vstack((X r2.T, target)).T
abalone lda = pd.DataFrame(data = lda data, columns = ['component1',
'component2','component3','component4','component5','label'] )
abalone temp lda = abalone lda.copy()
abalone temp 1da 2 = abalone temp 1da.drop(columns = ['label'])
abalone lda = (abalone temp lda 2 - abalone temp lda 2.min()) / (
abalone temp 1da 2.max() - abalone temp 1da 2.min())
abalone lda['label'] = abalone temp lda['label']
abalone lda['label'] = abalone lda['label'].astype('int')
abalone lda['label'] = abalone lda['label'].astype('category')
#print(abalone lda.head())
```

```
X_train_lda, X_test_lda, Y_train_lda, Y_test_lda = train_test_split(data_lda,
target_lda, test_size=0.2 , random_state=27)
```

# **Naive Bayes Classifier**

5-Fold CV for Naive Bayes & KNN classifier

```
In [64]:
        kfold = KFold(n splits=5)
        clrs = []
        clrs.append(KNeighborsClassifier(n neighbors=23, metric = 'manhattan',
        weights = "distance"))
        clrs.append(GaussianNB())
        clrs.append(MultinomialNB())
        clrs.append(ComplementNB())
        cv results = []
        cv results pca = []
        cv results lda = []
        for clr in clrs :
             cv results.append(cross val score(clr, data, target , scoring =
        'accuracy', cv = kfold, n jobs=-1))
            cv results pca.append(cross val score(clr, data pca, target pca, scoring
        = 'accuracy', cv = kfold, n jobs=-1))
            cv results lda.append(cross val score(clr, data lda, target lda, scoring
        = 'accuracy', cv = kfold, n jobs=-1))
        cv means = []
        cv means pca = []
        cv_means_lda = []
        for cv result in cv results:
            cv means.append(cv result.mean())
        for cv result in cv results pca:
            cv means pca.append(cv result.mean())
        for cv result in cv results lda:
            cv means lda.append(cv result.mean())
        cv df = pd.DataFrame({"Mean Accuracy":cv means, "Algo":
        ['KNeighborsClassifier','GaussianNB','MultinomialNB','ComplementNB']})
        cv df pca = pd.DataFrame({"Mean Accuracy":cv means pca,"Algo":
```

```
['KNeighborsClassifier','GaussianNB','MultinomialNB','ComplementNB']})
cv_df_lda = pd.DataFrame({"Mean_Accuracy":cv_means_lda,"Algo":
['KNeighborsClassifier','GaussianNB','MultinomialNB','ComplementNB']})

#print(cv_df)
#print(cv_df_pca)
#print(cv_df_lda)
```

```
abalone raw abalone pca abalone lda
Algo
KNeighborsClassifier 0.250907
                               0.249947
                                           0.253777
                     0.230326
                               0.232716
GaussianNB
                                          0.251631
MultinomialNB
                    0.206377
                               0.165673
                                          0.164955
ComplementNB
                     0.170705
                               0.171898
                                          0.228406
```

#### **Comments**

• Complement Naive Bayes performs poorly, the datset is unbalanced but due to multiclass classification it does not perform as expected for unbalanced datsets.

### **Decision Trees Classifier**

### 5-Fold CV and hyperparameter tuning

```
In [45]: dtc = DecisionTreeClassifier()
   dtc.fit(X_train, Y_train)
   Y_pred = dtc.predict(X_test)
   #cm=confusion_matrix(Y_test, Y_pred)
   #print(confusion_matrix(Y_test, Y_pred))
   #print(classification_report(Y_test, Y_pred))
   print("Accuracy Score for Decision Tree classifier without hyper parameter
   tuning is: ",accuracy_score(Y_test, Y_pred))
   #disp =
   ConfusionMatrixDisplay(confusion_matrix=cm,display_labels=dtc.classes_)
   #disp.plot()

grid_params = {
    #'criterion' : ['gini', 'entropy'],
    'max_depth' : range(2,25,1),
```

```
'min samples split' : range(2, 5, 1),
     'min samples leaf' : range(2, 5, 1)
grid search = GridSearchCV(dtc, grid params, cv = 5, n jobs = -1, verbose =
1)
grid search.fit(X train, Y train)
dtc = grid search.best estimator
Y pred = dtc.predict(X test)
print("Accuracy Score is :",accuracy score(Y test, Y pred))
 # best parameters and best score
print("Best Parameters are : ", grid search.best params )
#print("Best Score is : ",grid search.best score )
dt raw acc = accuracy score(Y test, Y pred)
dt raw set=grid search.best params
grid search.fit(X train pca, Y train pca)
dtc = grid search.best estimator
Y pred pca = dtc.predict(X test pca)
print("Accuracy Score using PCA is :",accuracy score(Y test pca, Y pred pca))
 # best parameters and best score
print("Best Parameters using PCA are : ", grid search.best params )
#print("Best Score using PCA is : ",grid_search.best_score_)
dt pca acc = accuracy score(Y test pca, Y pred pca)
dt pca set=grid search.best params
grid search.fit (X train lda, Y train lda)
dtc = grid search.best estimator
Y pred lda = dtc.predict(X test lda)
print("Accuracy Score using LDA is :",accuracy_score(Y_test_lda, Y_pred_lda))
 # best parameters and best score
print("Best Parameters using LDA are : ", grid search.best params )
 #print("Best Score using LDA is : ",grid search.best score )
dt lda acc = accuracy score(Y test lda, Y pred lda)
dt lda set=grid search.best params
Accuracy Score for Decision Tree classifier without hyper parameter tuning is: 0.19617224
```

```
880382775

Fitting 5 folds for each of 207 candidates, totalling 1035 fits

Accuracy Score is: 0.2535885167464115

Best Parameters are: {'max_depth': 4, 'min_samples_leaf': 2, 'min_samples_split': 2}

Fitting 5 folds for each of 207 candidates, totalling 1035 fits

Accuracy Score using PCA is: 0.26435406698564595

Best Parameters using PCA are: {'max_depth': 4, 'min_samples_leaf': 4, 'min_samples_split': 2}

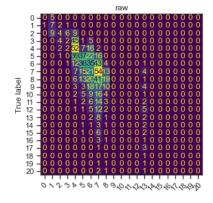
Fitting 5 folds for each of 207 candidates, totalling 1035 fits

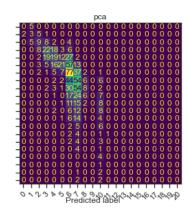
Accuracy Score using LDA is: 0.23444976076555024
```

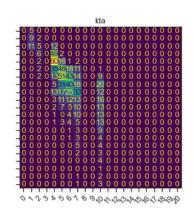
Best Parameters using LDA are: {'max depth': 3, 'min samples leaf': 2, 'min samples spli

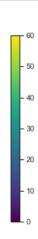
```
cm=confusion matrix(Y test, Y pred)
cm pca=confusion matrix(Y test pca, Y pred pca)
cm lda=confusion matrix(Y test_lda, Y_pred_lda)
cm dict = {
    "raw": cm,
    "pca": cm pca,
    "lda": cm lda
f, axes = plt.subplots(1, 3, figsize=(20, 5), sharey='row')
for i, (c,cm) in enumerate(cm dict.items()):
   #print(i,cm)
    disp = ConfusionMatrixDisplay(cm)
   disp.plot(ax=axes[i], xticks rotation=45)
   disp.ax .set title(c)
   disp.im .colorbar.remove()
   disp.ax .set xlabel('')
    if i!=0:
        disp.ax .set ylabel('')
f.text(0.4, 0.1, 'Predicted label', ha='left')
plt.subplots adjust(wspace=0.40, hspace=0.1)
f.suptitle('Confusion matrix', fontsize=12)
f.colorbar(disp.im , ax=axes)
plt.show()
```











plt.show()

```
In [48]:
        def plot acc vs depth(dataset):
             \max depth range = list(range(1, 25))
             acc list = []
             acc list pca = []
             acc list lda = []
             for depth in max depth range:
                 dtc w = DecisionTreeClassifier(max depth = depth)
                 if dataset[0] == 'raw':
                     dtc w.fit(X train, Y train)
                     score = dtc w.score(X test, Y test)
                     acc list.append(score)
                 if dataset[1] == 'pca':
                     dtc w.fit(X train pca, Y train pca)
                     score = dtc w.score(X test pca, Y test pca)
                     acc list pca.append(score)
                 if dataset[2] == 'lda':
                     dtc w.fit(X train lda, Y train lda)
                     score =dtc w.score(X test lda, Y test lda)
                     acc list lda.append(score)
             return (acc list, acc list pca, acc list lda)
        acc list,acc list pca,acc list lda = plot acc vs depth(['raw','pca','lda'])
        fig = plt.figure(figsize=(7,5))
        plt.plot(range(1,25),acc list, color = 'blue', marker ='o', mew
        =4, mfc='yellow', ls ='--' , linewidth=1.5, label='Accuracy Score raw')
        plt.plot(range(1,25),acc list pca, color = 'cyan',marker ='o',mew
        =4, mfc='yellow', ls ='--' , linewidth=1.5, label='Accuracy Score pca')
        plt.plot(range(1,25),acc list lda, color = 'm', marker = 'o', mew
        =4, mfc='yellow', ls ='--' , linewidth=1.5, label='Accuracy Score lda')
        plt.ylabel('Accuracy', fontsize = 15)
        plt.xlabel('Max Depth', fontsize = 15)
        plt.title('Accuracy vs Max Depth for wine dataset', fontsize = 18)
        plt.legend(loc='best', fontsize = 15)
        plt.tight layout()
```

# O.26 - Accuracy Score\_raw - Accuracy Score\_pca - Accuracy Score\_lda

10

# Splitting rules used for the trees

5

0.18

```
In [49]: dtc_g = DecisionTreeClassifier(criterion= 'entropy', max_depth=4,
    min_samples_leaf= 2, min_samples_split=2)
    dtc_g.fit(X_train,Y_train)
    fn=X_train.columns
    cn = abalone_raw.Rings.unique().astype(str)
    dot_data = tree.export_graphviz(dtc_g,
    filled=True, feature_names=fn, class_names=cn , rounded=True,
    special_characters=True)
    graph = graphviz.Source(dot_data)
    graph
```

15

Max Depth

20

```
Out[49]:
```

```
In [60]: from sklearn.tree import export_text
  dtc_g = DecisionTreeClassifier(criterion= 'entropy', max_depth=4,
  min_samples_leaf= 2, min_samples_split=2)
  dt_rules = dtc_g.fit(X_train,Y_train)
  fn_t=X_train.columns
  #print(type(fn_t.tolist()))
  tree_rules = export_text(dt_rules,feature_names = fn_t.tolist())
  print(tree_rules)
```

```
|--- class: 3
            |--- Shucked weight > 0.00
                |--- class: 4
        |--- Shell weight > 0.02
            |--- Shucked weight <= 0.04
                |--- class: 5
            |--- Shucked weight > 0.04
                |--- class: 6
    \mid--- Shell weight > 0.05
        |--- Sex <= 1.50
            |--- Shucked weight <= 0.16
                |--- class: 9
            |--- Shucked weight > 0.16
            | |--- class: 8
         --- Sex > 1.50
            \mid--- Shell weight \leq 0.12
                |--- class: 7
            \mid --- Shell weight > 0.12
               |--- class: 8
            |--- Shell weight > 0.15
    |--- Shell weight <= 0.29
        |--- Shucked weight <= 0.26
            |--- Shell weight <= 0.19
              |--- class: 8
            \mid --- Shell weight > 0.19
                |--- class: 10
        |--- Shucked weight > 0.26
            \mid--- Shell weight \leq 0.26
            | |--- class: 9
            |--- Shell weight > 0.26
               |--- class: 9
    |--- Shell weight > 0.29
        |--- Shucked weight <= 0.30
            |--- Height <= 0.17
              |--- class: 10
            |--- Height > 0.17
                |--- class: 13
        |--- Shucked weight > 0.30
            |--- Shell weight <= 0.41
               |--- class: 10
            |--- Shell weight > 0.41
           | |--- class: 11
```

#### **Comments**

• As observed in Assignment-1, weight columns were positively correlated with many other columns(namely; Height, Length, Diameter) and hence it is chosen as the root node for the split. Weight and Sex are the major contributors to splitting rules indicating that these are more important for classication.

#### Splitting rules used for the trees but fitted on page

```
In [51]: dtc_g = DecisionTreeClassifier(criterion= 'entropy', max_depth=4,
    min_samples_leaf= 2, min_samples_split=5)
    dtc_g.fit(X_train,Y_train)
    fn=X_train.columns
```

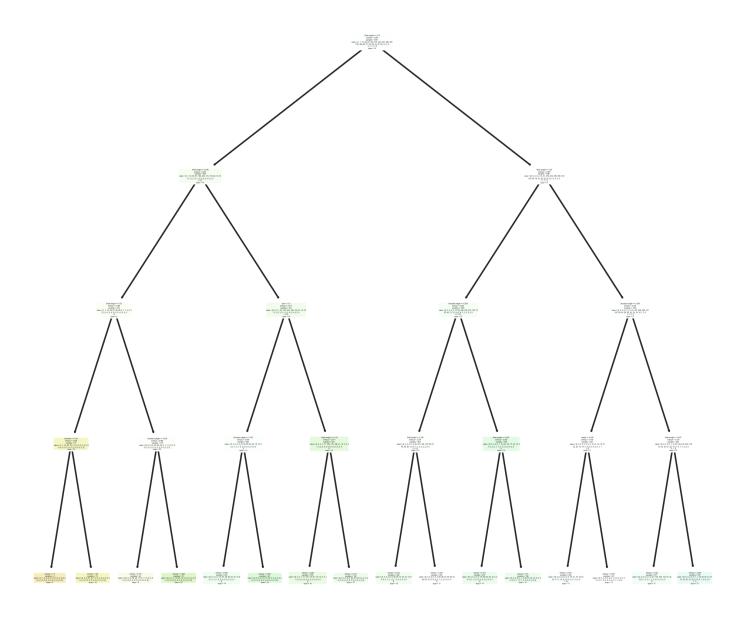
```
cn = abalone raw.Rings.unique().astype(str)
        fig, axes = plt.subplots(nrows = 1, ncols = 1, figsize = (8,8), dpi=300)
        tree.plot tree(dtc q, filled=True, feature names=fn, class names=cn
        , rounded=True)
        #fig.savefig('dtc tree plot 3.png')
Out[51]: [Text(930.0, 1630.8, 'Shell weight <= 0.15\nentropy = 3.593\nsamples = 3341\nvalue = [1,
       1, 10, 46, 87, 202, 310, 453, 557, 520, 387\n212, 166, 98, 77, 54, 50, 33, 27, 22, 13, 3,
       5\n2, 1, 1, 2, 1]\nclass = 14'),
       Text(465.0, 1268.4, 'Shell weight <= 0.046\nentropy = 2.972\nsamples = 980\nvalue = [1,
       1, 10, 46, 87, 189, 249, 174, 104, 62, 19, 18\n11, 3, 2, 3, 0, 1, 0, 0, 0, 0, 0, 0, 0\n
       0, 0] \ln s = 16'),
       Text(232.5, 906.0, 'Shell weight <= 0.02\nentropy = 2.386\nsamples = 219\nvalue = [1, 1,
       10, 45, 67, 59, 26, 8, 1, 1, 0, 0, 0\n0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0\n0]\nclass
       = 8'),
       Text(116.25, 543.600000000001, 'Diameter <= 0.08\nentropy = 2.028\nsamples = 74\nvalue =
       lass = 10'),
       Text(58.125, 181.200000000000000, 'entropy = 1.5\nsamples = 4\nvalue = [1, 1, 2, 0, 0, 0, 0]
       Text(174.375, 181.20000000000005, 'entropy = 1.84\nsamples = 70\nvalue = [0, 0, 8, 35, 1]
       Text(348.75, 543.6000000000001, 'Shucked weight <= 0.036\nentropy = 2.085\nsamples = 145
       0, 0, 0] \land class = 20'),
       Text(290.625, 181.2000000000005, 'entropy = 2.179\nsamples = 94\nvalue = [0, 0, 0, 9, 3
       6, 28, 11, 8, 1, 1, 0, 0, 0, 0 \setminus 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0] \nclass = 8'),
       Text(406.875, 181.20000000000000, 'entropy = 1.628\nsamples = 51\nvalue = [0, 0, 0, 1, 1]
       Text(697.5, 906.0, 'Sex \leq 1.5\nentropy = 2.714\nsamples = 761\nvalue = [0, 0, 0, 1, 20,
       130, 223, 166, 103, 61, 19, 18 \ln 1, 3, 2, 3, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0 \ln 0, 0 \ln 0
       16'),
       Text (581.25, 543.600000000001, 'Shucked weight \leq 0.156 \cdot \text{nentropy} = 2.905 \cdot \text{nsamples} = 280
       0, 0, 0 \le 0 \le 14'),
       Text(523.125, 181.20000000000000, 'entropy = 2.952\nsamples = 234\nvalue = [0, 0, 0, 0,
       3, 15, 34, 38, 59, 43, 15, 13, 8 \cdot n2, 2, 1, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0 \cdot n0] \cdot nclass = 1
       4'),
       Text(639.375, 181.20000000000000, 'entropy = 1.942\nsamples = 46\nvalue = [0, 0, 0, 0, 0, 0, 0]
       Text(813.75, 543.600000000001, 'Shell weight \leq 0.118\nentropy = 2.371\nsamples = 481\nv
       0, 0, 0 \le 0 \le 16'),
       Text(755.625, 181.20000000000005, 'entropy = 2.266\nsamples = 355\nvalue = [0, 0, 0, 1, 1]
       7, 101, 137, 61, 19, 10, 4, 3, 1\n1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0\n0]\nclass = 1
       6'),
       Text(871.875, 181.2000000000005, 'entropy = 2.214\nsamples = 126\nvalue = [0, 0, 0, 0, 0]
       0, 8, 38, 47, 22, 5, 0, 2, 2, 0 \setminus n0, 2, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0] \setminus nclass = 19'),
       Text(1395.0, 1268.4, 'Shell weight \leq 0.29 \cdot \text{nentropy} = 3.336 \cdot \text{nsamples} = 2361 \cdot \text{nvalue} = [0, 1395.0]
       0, 0, 0, 0, 13, 61, 279, 453, 458, 368, 194 n 155, 95, 75, 51, 50, 32, 27, 22, 13, 3, 5, 2
       n1, 1, 2, 1] nclass = 11'),
       Text(1162.5, 906.0, 'Shucked weight <= 0.256\nentropy = 3.083\nsamples = 1206\nvalue =
       [0, 0, 0, 0, 0, 10, 58, 246, 300, 212, 129, 73 \n52, 40, 31, 15, 14, 8, 8, 3, 3, 2, 2, 0, 0]
       n0, 0, 0] nclass = 14'),
       Text(1046.25, 543.600000000001, 'Shell weight \leq 0.192 \neq 3.276 = 801 = 801
```

 $2, 0, 0 \neq 0, 0 \neq 0, 0 \neq 0$ 

```
Text(988.125, 181.200000000000000, 'entropy = 2.912\nsamples = 325\nvalue = [0, 0, 0, 0, 0, 0]
0, 5, 35, 93, 73, 44, 30, 13, 8 \ 8, 3, 1, 2, 1, 0, 1, 0, 0, 0, 0, 0, 0 \ 0] \ class = 1
9'),
 Text(1104.375, 181.2000000000000, 'entropy = 3.339\nsamples = 476\nvalue = [0, 0, 0, 0,
11'),
 Text(1278.75, 543.600000000001, 'Shell weight \leq 0.257 \neq 2.525 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 405 = 
value = [0, 0, 0, 0, 0, 1, 19, 93, 141, 77, 44, 12, 3\n4, 3, 2, 3, 1, 1, 0, 1, 0, 0, 0, 0,
0, 0 \neq 0 
 Text(1220.625, 181.2000000000000, 'entropy = 2.104\nsamples = 207\nvalue = [0, 0, 0, 0,
4'),
 Text(1336.875, 181.20000000000005, 'entropy = 2.73\nsamples = 198\nvalue = [0, 0, 0, 0, 0]
0, 1, 4, 28, 61, 45, 34, 8, 3, 4 \ 3, 2, 2, 1, 1, 0, 1, 0, 0, 0, 0, 0, 0, 0] \ class = 14'),
 Text(1627.5, 906.0, 'Shucked weight <= 0.298\nentropy = 3.335\nsamples = 1155\nvalue =
[0, 0, 0, 0, 0, 3, 3, 33, 153, 246, 239, 121 \n103, 55, 44, 36, 36, 24, 19, 19, 10, 1, 3, 2]
n1, 1, 2, 1] nclass = 11'),
 Text(1511.25, 543.6000000000001, 'Height \leq 0.166\nentropy = 3.758\nsamples = 227\nvalue
= [0, 0, 0, 0, 0, 2, 0, 5, 15, 31, 19, 19, 31\n22, 20, 15, 14, 7, 10, 8, 3, 0, 3, 0, 1, 1
n1, 0]nclass = 11'),
 Text(1453.125, 181.2000000000000, 'entropy = 3.672\nsamples = 202\nvalue = [0, 0, 0, 0,
0, 2, 0, 4, 15, 31, 19, 19, 27 \land 11, 10, 6, 10, 8, 3, 0, 1, 0, 0, 0 \land 1, 0] \land 12 = 0
11'),
 Text(1569.375, 181.20000000000005, 'entropy = 3.094\nsamples = 25\nvalue = [0, 0, 0, 0, 0]
0, 0, 0, 1, 0, 0, 0, 0, 4, 4 \ 1, 0, 0, 0, 0, 0, 0, 1, 1, 0, 0] \ = 13'),
 Text(1743.75, 543.600000000001, 'Shell weight \leq 0.407 \neq 3.126 = 928 \neq 3.126 \neq 
value = [0, 0, 0, 0, 0, 1, 3, 28, 138, 215, 220, 102\n72, 33, 24, 21, 22, 17, 9, 11, 7, 1,
0, 2, 0 \neq 0, 1, 1] \neq 12'
 Text(1685.625, 181.2000000000000, 'entropy = 2.702\nsamples = 571\nvalue = [0, 0, 0, 0,
0, 1, 2, 27, 120, 165, 133, 51, 35 \ n15, 8, 1, 4, 5, 1, 1, 2, 0, 0, 0, 0, 0, 0 \ n0] \ nclass = 0
11'),
 Text(1801.875, 181.2000000000000, 'entropy = 3.398\nsamples = 357\nvalue = [0, 0, 0, 0,
```

 $0, 0, 1, 1, 18, 50, 87, 51, 37 \ln 8, 16, 20, 18, 12, 8, 10, 5, 1, 0, 2, 0, 0 \ln 1, 1] \ln 2$ 

= 12')]



# **Random Forest Classifier**

# 5-Fold CV and hyperpararmeter tuning

```
grid params = {
    'n estimators': [2,5,200, 500],
    'criterion' : ['entropy'],
    'max depth' : range(2,25,1),
    #'max features': range(3,10,1),
    #'min samples split' : range(2, 10, 1),
    #'min samples leaf' : range(2, 10, 1)
grid search = GridSearchCV(rfc, grid params, cv = 5, n jobs = -1, verbose =
#print(grid search.get params().keys())
grid search.fit(X train, Y train)
rfc = grid search.best estimator
Y pred = rfc.predict(X test)
print("Accuracy Score is :",accuracy score(Y test, Y pred))
rf raw acc=accuracy score(Y test, Y pred)
# best parameters and best score
print("Best Parameters are : ", grid search.best params )
rf raw set=grid search.best params
#print("Best Score is : ", grid search.best score )
ac df=pd.DataFrame(grid_search.cv_results_['params'])
#Creating a data frame with hyperparameters and accuracy
ac df["accuracy"]=grid search.cv results ['mean test score']
grid search.fit(X train pca, Y train pca)
rfc pca = grid search.best estimator
Y pred pca = rfc pca.predict(X test pca)
print("Accuracy Score using PCA is :",accuracy score(Y test pca, Y pred pca))
rf pca acc=accuracy score(Y test pca, Y pred pca)
# best parameters and best score
print("Best Parameters using PCA are : ", grid search.best params )
rf pca set=grid search.best params
#print("Best Score using PCA is : ",grid search.best score )
ac df pca=pd.DataFrame(grid search.cv results ['params'])
#Creating a data frame with hyperparameters and accuracy
ac df pca["accuracy"]=grid search.cv results ['mean test score']
grid search.fit(X train lda, Y train lda)
rfc lda = grid search.best estimator
Y pred lda = rfc lda.predict(X test lda)
print("Accuracy Score using LDA is :",accuracy score(Y test lda, Y pred lda))
rf lda acc=accuracy score(Y test lda, Y pred lda)
```

```
# best parameters and best score
print("Best Parameters using LDA are : ",grid_search.best_params_)
#print("Best Score using LDA is : ",grid_search.best_score_)
rf_lda_set=grid_search.best_params_
ac_df_lda=pd.DataFrame(grid_search.cv_results_['params'])
#Creating a data frame with hyperparameters and accuracy
ac_df_lda["accuracy"]=grid_search.cv_results_['mean_test_score']
```

```
Accuracy Score for Forest classifier without hyper parameter tuning is: 0.246411483253588 53

Fitting 5 folds for each of 92 candidates, totalling 460 fits
Accuracy Score is: 0.27392344497607657

Best Parameters are: {'criterion': 'entropy', 'max_depth': 5, 'n_estimators': 5}

Fitting 5 folds for each of 92 candidates, totalling 460 fits
Accuracy Score using PCA is: 0.2679425837320574

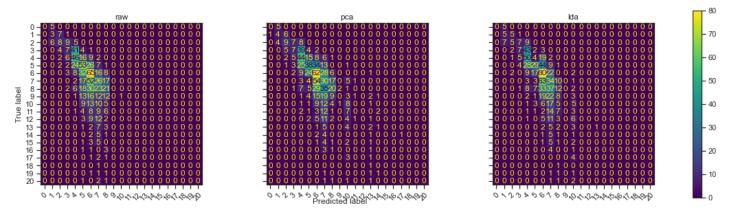
Best Parameters using PCA are: {'criterion': 'entropy', 'max_depth': 6, 'n_estimators': 200}

Fitting 5 folds for each of 92 candidates, totalling 460 fits
Accuracy Score using LDA is: 0.27272727272727

Best Parameters using LDA are: {'criterion': 'entropy', 'max_depth': 4, 'n_estimators': 500}
```

```
In [53]:
        cm=confusion matrix(Y test, Y pred)
        cm pca=confusion matrix(Y test pca, Y pred pca)
        cm lda=confusion matrix(Y test lda, Y pred lda)
         cm dict = {
            "raw": cm,
            "pca": cm pca,
            "lda": cm lda
        f, axes = plt.subplots(1, 3, figsize=(20, 5), sharey='row')
        for i, (c,cm) in enumerate(cm dict.items()):
             #print(i,cm)
             disp = ConfusionMatrixDisplay(cm)
             disp.plot(ax=axes[i], xticks rotation=45)
            disp.ax .set title(c)
             disp.im .colorbar.remove()
            disp.ax .set xlabel('')
             if i!=0:
                 disp.ax .set ylabel('')
        f.text(0.4, 0.1, 'Predicted label', ha='left')
        plt.subplots adjust(wspace=0.40, hspace=0.1)
```

```
f.colorbar(disp.im_, ax=axes)
plt.show()
```



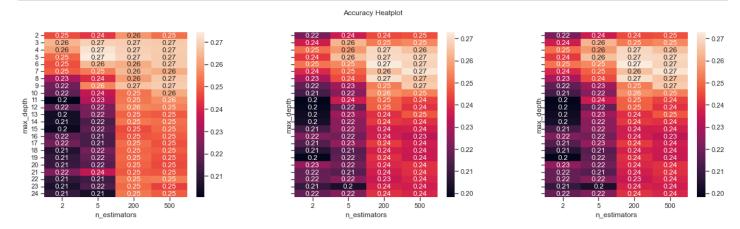
#### Heatmap of max\_depth vs n\_estimators

```
In [54]: #Pivoting the dataframe for plotting heat map
    ac_df=ac_df.pivot(index='max_depth',columns='n_estimators',values='accuracy')
    ac_df_pca=ac_df_pca.pivot(index='max_depth',columns='n_estimators',values='accuracy')
    ac_df_lda=ac_df_lda.pivot(index='max_depth',columns='n_estimators',values='accuracy')

#Plotting the graph
fig, ax =plt.subplots(1,3,figsize=(20, 5), sharey='row')

sns.heatmap(ac_df,annot=True, ax=ax[0])
sns.heatmap(ac_df_pca,annot=True, ax=ax[1])
sns.heatmap(ac_df_pca,annot=True, ax=ax[2])

plt.subplots_adjust(wspace=0.40, hspace=0.1)
fig.suptitle('Accuracy Heatplot', fontsize=12)
plt.show()
```



# **Gradient Boosting Classifier**

#### 5-Fold CV and hyperparameter tuning

```
gbc = GradientBoostingClassifier()
gbc.fit(X train, Y train)
Y pred = gbc.predict(X test)
cm=confusion matrix(Y test, Y pred)
#print(confusion matrix(Y test, Y pred))
#print(classification report(Y test, Y pred))
print ("Accuracy Score for Gradient Boosting classifier without hyper
parameter tuning is: ",accuracy_score(Y_test, Y_pred))
#disp =
ConfusionMatrixDisplay(confusion matrix=cm, display labels=dtc.classes)
#disp.plot()
grid params = {
    'n estimators': [2,120,200,500,800],
    'learning rate' : [0.01, 0.1,1],
    #'criterion' : [None,'mse','mae'],
    #'max depth' : range(2,22,2),
    #'max features': range(3,10,1),
    #'min samples split' : range(2, 10, 1),
    #'min samples leaf' : range(2, 10, 1)
grid search = GridSearchCV(gbc, grid params, cv = 5, n jobs = -1, verbose =
True)
#print(grid search.get params().keys())
grid search.fit(X train, Y train)
gbc = grid search.best estimator
Y pred = gbc.predict(X test)
print("Accuracy Score is :",accuracy score(Y test, Y pred))
gbc acc=accuracy score(Y test, Y pred)
# best parameters and best score
print("Best Parameters are : ", grid search.best params )
gbc set=grid search.best params
#print("Best Score using PCA is : ",grid search.best score )
gbc ac df=pd.DataFrame(grid search.cv results ['params'])
#Creating a data frame with hyperparameters and accuracy
gbc ac df["accuracy"]=grid search.cv results ['mean test score']
grid search.fit(X train pca, Y train pca)
gbc = grid search.best estimator
Y pred pca = gbc.predict(X test pca)
print("Accuracy Score is :",accuracy score(Y test pca, Y pred pca))
```

```
gbc pca acc=accuracy score(Y test pca, Y pred pca)
# best parameters and best score
print("Best Parameters are : ", grid search.best params )
gbc_pca_set=grid_search.best params
#print("Best Score using PCA is : ",grid search.best score )
gbc ac df pca=pd.DataFrame(grid search.cv results ['params'])
 #Creating a data frame with hyperparameters and accuracy
gbc ac df pca["accuracy"]=grid search.cv results ['mean test score']
grid search.fit(X train lda, Y train lda)
gbc = grid search.best estimator
Y pred lda = gbc.predict(X test lda)
print("Accuracy Score is :",accuracy score(Y test lda, Y pred lda))
gbc lda acc=accuracy score(Y test lda, Y pred lda)
# best parameters and best score
print("Best Parameters are : ", grid search.best params )
gbc lda set=grid search.best params
#print("Best Score using PCA is : ",grid search.best score )
gbc ac df lda=pd.DataFrame(grid search.cv results ['params'])
 #Creating a data frame with hyperparameters and accuracy
gbc ac df lda["accuracy"]=grid search.cv results ['mean test score']
Accuracy Score for Gradient Boosting classifier without hyper parameter tuning is: 0.2535
885167464115
```

```
885167464115

Fitting 5 folds for each of 15 candidates, totalling 75 fits

Accuracy Score is: 0.26913875598086123

Best Parameters are: {'learning_rate': 0.01, 'n_estimators': 120}

Fitting 5 folds for each of 15 candidates, totalling 75 fits

Accuracy Score is: 0.26435406698564595

Best Parameters are: {'learning_rate': 0.01, 'n_estimators': 200}

Fitting 5 folds for each of 15 candidates, totalling 75 fits

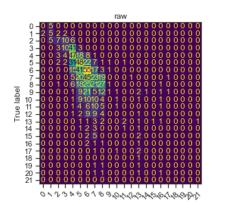
Accuracy Score is: 0.28708133971291866

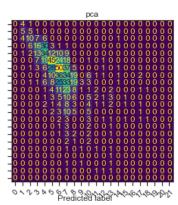
Best Parameters are: {'learning rate': 0.01, 'n estimators': 120}
```

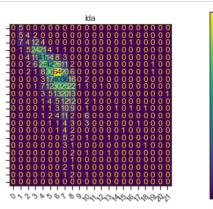
```
In [56]:
    cm=confusion_matrix(Y_test, Y_pred)
    cm_pca=confusion_matrix(Y_test_pca, Y_pred_pca)
    cm_lda=confusion_matrix(Y_test_lda, Y_pred_lda)
    cm_dict = {
        "raw": cm,
        "pca": cm_pca,
        "lda": cm_lda
    }

    f, axes = plt.subplots(1, 3, figsize=(20, 5), sharey='row')
    for i,(c,cm) in enumerate(cm_dict.items()):
        #print(i,cm)
```

```
disp = ConfusionMatrixDisplay(cm)
    disp.plot(ax=axes[i], xticks rotation=45)
    disp.ax .set title(c)
    disp.im .colorbar.remove()
    disp.ax .set xlabel('')
    if i!=0:
        disp.ax .set ylabel('')
f.text(0.4, 0.1, 'Predicted label', ha='left')
plt.subplots adjust(wspace=0.40, hspace=0.1)
f.colorbar(disp.im , ax=axes)
plt.show()
```

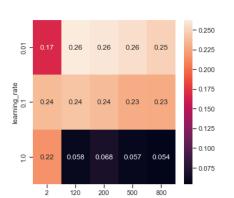


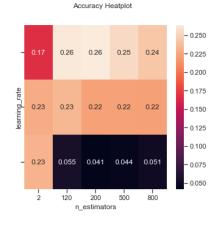


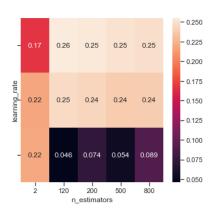


# Heatplot of learning\_rate and n\_estimators

```
In [57]:
        #Pivoting the dataframe for plotting heat map
        gbc ac df=gbc ac df.pivot(index='learning rate',columns='n estimators',values=
        gbc ac df pca=gbc ac df pca.pivot(index='learning rate',columns='n estimators'
        gbc ac df lda=gbc ac df lda.pivot(index='learning rate',columns='n estimators'
        #Plotting the graph
        fig, ax =plt.subplots(1,3,figsize=(20, 5), sharey='row')
        sns.heatmap(gbc ac df,annot=True, ax=ax[0])
        sns.heatmap(gbc ac df pca,annot=True ,ax=ax[1])
        sns.heatmap(gbc ac df lda,annot=True ,ax=ax[2])
        plt.subplots adjust(wspace=0.40, hspace=0.1)
        fig.suptitle('Accuracy Heatplot', fontsize=12)
        plt.show()
```







#### **Final Results**

```
abalone_raw abalone_pca
                                                    abalone_lda
   KNeighborsClassifier
                            0.250907
                                          0.249947
                                                        0.253777
1
           GaussianNB
                            0.230326
                                          0.232716
                                                        0.251631
2
        MultinomialNB
                            0.206377
                                          0.165673
                                                        0.164955
       ComplementNB
                            0.170705
                                          0.171898
                                                        0.228406
3
                                          0.264354
4
          Decision Tree
                            0.253589
                                                        0.234450
                                          0.267943
5
        Random Forest
                            0.273923
                                                        0.272727
                            0.269139
                                          0.264354
                                                        0.287081
6
     Gradient Boosting
```

```
In [67]: #print(dt_raw_set.keys())
    #print(dt_raw_set.values(),dt_pca_set.values(),dt_lda_set.values())
    dt_s =[dt_raw_set,dt_pca_set,dt_lda_set]
    df_dt_settings = pd.DataFrame(dt_s)
    df_dt_settings['Algo'] = 'Decision Tree'
    df_dt_settings['Data'] = ['raw','pca','lda']
    display(df_dt_settings)
```

```
    max_depth
    min_samples_leaf
    min_samples_split
    Algo
    Data

    0
    4
    2
    2
    Decision Tree
    raw
```

```
max_depthmin_samples_leafmin_samples_splitAlgoData1442Decision Treepca2322Decision TreeIda
```

```
criterionmax_depthn_estimatorsAlgoData0entropy55Random Forestraw1entropy6200Random Forestpca2entropy4500Random ForestIda
```

```
In [69]: gbc_s =[gbc_set,gbc_pca_set,gbc_lda_set]
  df_gbc_settings = pd.DataFrame(gbc_s)
  df_gbc_settings['Algo'] = 'Gradient Boosting'
  df_gbc_settings['Data'] = ['raw','pca','lda']
  display(df_gbc_settings)
```

```
learning_raten_estimatorsAlgoData00.01120Gradient Boostingraw10.01200Gradient Boostingpca20.01120Gradient BoostingIda
```

```
In [2]: knn_setting = ({'n_neighbors':66 , 'metric': 'minkowski' , 'weights':
    "uniform",'p':2})
    gnb_settings = ("default")
    mnb_settings = ("default")
    cnb_settings = ("default")
    dt_settings =
        ({'criterion':'Entropy', 'max_depth':4, 'min_samples_leaf':4, 'min_samples_split':
        rf_settings = ({'criterion':'Entropy', 'max_depth':5, 'n_estimators':500})
        gb_settings = ({'learning_rate':0.01, 'n_estimators':120})
        settings_all =
        [knn_setting,gnb_settings,mnb_settings,cnb_settings,dt_settings,dt_settings,rf_settings_all
        df_final['settings'] =settings_all
```

In [5]:

Out[5]:

display(df final)

:	Algo	settings	abalone_raw	abalone_pca	abalone_lda
	<b>0</b> KNeighborsClassifier	{'n_neighbors':66 , 'metric': 'minkowski' , 'w	0.250907	0.249947	0.253777
	<b>1</b> GaussianNB	default	0.230326	0.232716	0.251631
	2 MultinomialNB	default	0.206377	0.165673	0.164955
	3 ComplementNB	default	0.170705	0.171898	0.228406
	4 Decision Tree	$ \label{lem:condition} \mbox{\cite{criterion':'Entropy','max\_depth':4,'min\_samp} } \\$	0.253589	0.264354	0.234450
	<b>5</b> Random Forest	{'criterion':'Entropy','max_depth':5,'n_estima	0.273923	0.267943	0.272727
	<b>6</b> Gradient Boosting	{'learning_rate':0.01,'n_estimators':120}	0.269139	0.264354	0.287081

#### **Comments**

- Gradient Boosting performed best with the pipeline (abalone -> scaling -> Ida -> scaling -> Gradient Boosting Classifier)
- Dimensionality Reduction helped in Decision tree, Random Forest and Gradient Boosting algorithms, while PCA gav econsistent accuracies with mentioned algorithms, LDA performed well only with Gradient Boosting and Random Forest algorithms.
- Decision tree and Random Forest gave better accuracy with lesser max depth as compared to wine dataset.

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randomforest#:~:text=There%20is%20no%20problem%20with,two%20features%2C%20Age%20and%20Sex%20.