```
import warnings
warnings.filterwarnings('ignore')
from sklearn.metrics import mean squared error, r2 score,
accuracy score
from sklearn.linear model import LinearRegression
import pandas as pd
import numpy as np
from plotnine import *
from sklearn.decomposition import PCA
from sklearn import metrics
from sklearn.linear model import LogisticRegression # Logistic
Regression Model
from sklearn.preprocessing import StandardScaler #Z-score variables
from sklearn.metrics import accuracy score, confusion matrix
from sklearn.metrics import accuracy_score, confusion_matrix,
plot confusion matrix
from sklearn.model selection import train test split # simple TT split
from sklearn.model selection import KFold # k-fold cv
from sklearn.model selection import LeaveOneOut #LOO cv
from sklearn.model selection import cross val score # cross validation
metrics
from sklearn.model selection import cross val predict # cross
validation metrics
from sklearn.cluster import DBSCAN
from sklearn.neighbors import NearestNeighbors
from sklearn.cluster import KMeans
from sklearn.mixture import GaussianMixture
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import silhouette score
rawData =
pd.read csv("https://gist.githubusercontent.com/armgilles/194bcff35001
e7eb53a2a8b441e8b2c6/raw/92200bc0a673d5ce2110aaad4544ed6c4010f687/
pokemon.csv")
rawData.head()
newData = rawData.copy()
df = newData.copy()
df['Legendary'] = df['Legendary'].astype(int)
df
                           Name
                                  Type 1 Type 2 Total HP Attack
Defense \
                      Bulbasaur
                                   Grass
                                          Poison
                                                    318 45
                                                                 49
0
       1
49
1
       2
                                                    405
                                                                 62
                        Ivysaur
                                   Grass
                                          Poison
                                                         60
63
2
       3
                                                                 82
                       Venusaur
                                   Grass Poison
                                                    525 80
83
3
       3 VenusaurMega Venusaur
                                   Grass Poison
                                                    625 80
                                                                100
123
```

4 43	4	Charmander				Fire	NaN	309	39	52
795 150 796 110 797 60 798 60 799 120	719	Diancie				Rock	Fairy	600	50	100
	719	DiancieMega Diancie				Rock	Fairy	700	50	160
	720	HoopaHoopa Confined				Psychic	Ghost	600	80	110
	720	HoopaHoopa Unbound				Psychic	Dark	680	80	160
	721			Volcani	on	Fire	Water	600	80	110
0 1 2 3 4 	Sp.	Atk Sp 65 80 100 122 60 	p. Def 65 80 100 120 50	Speed 45 60 80 80 65	Gei	neration 1 1 1 1 1	Legenda	y 0 0 0 0 0		
796		160	110	110		6		1		
797		150	130	70		6		1		
798 799		170 130	130 90	80 70		6 6		1 1		
[800 rows x 13 columns]										

[800 rows x 13 columns]

1)

When creating a Logistic Regression model is it best to use HP, Attack, Defense, and Speed or Special Attack and Special Defense as predictors to determine if a Pokemon is Legendary?

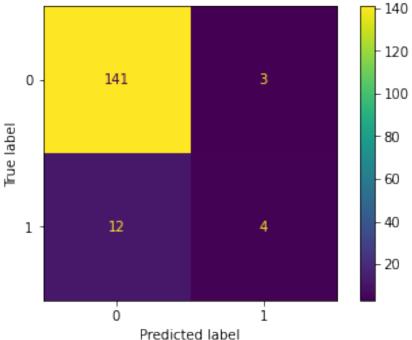
The accuracy scores of the models were 90.63% and 91.88% showing that both models were very effective. However in the future I cn create a better model for HP, Attack, Defense, and Speed if I dont use Attack as a predictor because when graphing the coefficients you can see that Attack has the lowest coefficient of 0.46 showing us that it has the lowest impact on determining if a Pokemon is Legendary.

However, I would definitely go with my second model to make prediction because with only 2 predictors it was just as accurate as the first model.

This model differed from my analysis to determine which predictor groups were the most acurate rather than which specific predictors.

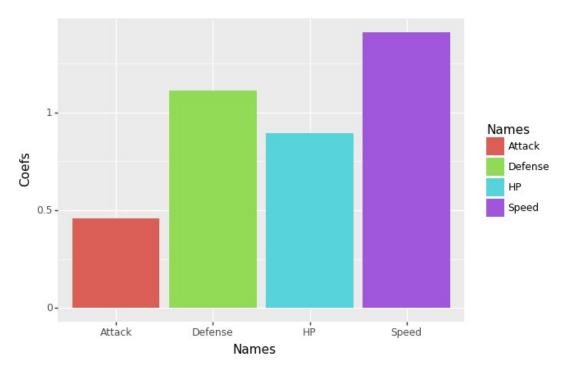
```
predictors = ["HP", "Attack", "Defense", "Speed"]
X_train, X_test, y_train, y_test = train_test_split(df[predictors],
```

```
df["Legendary"], test_size=0.2)
z = StandardScaler()
z.fit(X_train[predictors])
X_test[predictors]=z.transform(X_test[predictors])
X_train[predictors]=z.transform(X_train[predictors])
lr = LogisticRegression()
lr.fit(X_train, y_train)
y_pred = lr.predict(X_train)
confusion_matrix(y_train, y_pred)
X = df[predictors]
y = df["Legendary"]
plot_confusion_matrix(lr,X_test,y_test)
<sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at
0x7f5d372d6190>
```

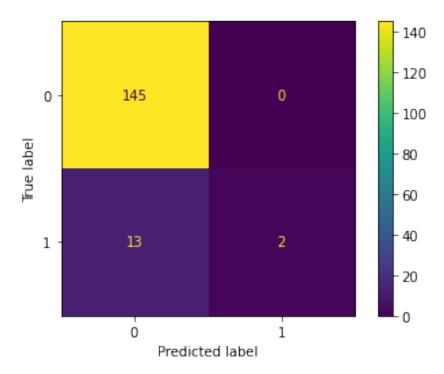


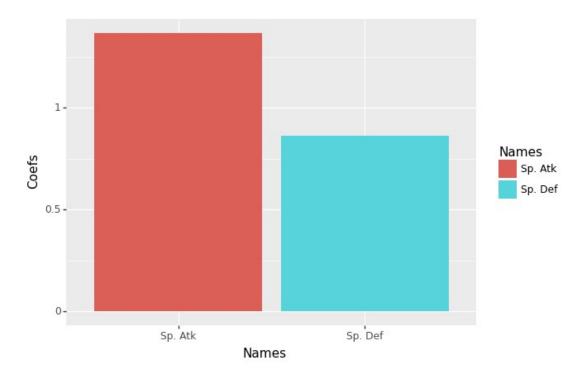
```
print("Accuracy score:",accuracy_score(y_test, lr.predict(X_test)))
Accuracy score: 0.90625
coef = pd.DataFrame({"Coefs": lr.coef_[0],
                     "Names": predictors})
coef
      Coefs
               Names
  0.893368
                  HP
1
  0.459688
              Attack
  1.112682
             Defense
  1.408566
               Speed
```

```
(ggplot(coef, aes(x = "Names", y = "Coefs", fill = "Names" )) +
geom bar(stat = "identity"))
```



<ggplot: (8752395868793)> predictors2 = ["Sp. Atk", "Sp. Def"] X_train, X_test, y_train, y_test = train_test_split(df[predictors2], df["Legendary"], test size=0.2) z = StandardScaler() z.fit(X train[predictors2]) X test[predictors2]=z.transform(X test[predictors2]) X train[predictors2]=z.transform(\overline{X} train[predictors2]) lr = LogisticRegression() lr.fit(X train, y train) y pred = lr.predict(X train) confusion_matrix(y_train, y_pred) X = df[predictors2]y = df["Legendary"] plot_confusion_matrix(lr,X_test,y_test) <sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at</pre> 0x7f5d37103ad0>





<ggplot: (8752395855841)>

2)

When using Total and Sp.Attack (the most influential predictor for Legendaries) to cluster Legendaries what is the best clustering model to use?

When creating clustering models I determined that the best model to use was KNN because KNN ended up having an accuracy score of 93.13% which could be because my model was very accurate in counting how many neighbors the different Pokemon had to classify them.

DBSCAN was very inaccurate because the model was very random and did not have too much noise, so I was not surprised to see that the silhouette score was a low 0.46, showing that the model clustered badly.

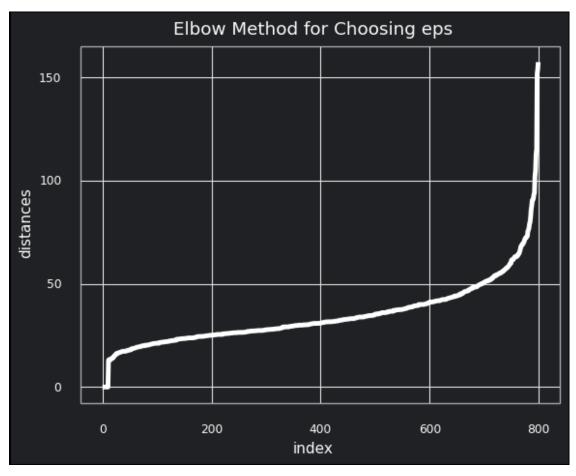
Gaussian Mixture method was the most inefficient, possibly because the data was so random, and had a lower silhouette score of 0.175.

Here I used DBSCAN to cluster Legendaries, and ended up realizing that the randomness of the data and lack of noise points were really bad for my model.

In my analysis I wantd to use DBSCAN to cluster types, but I decided to cluster legendaries for accuracy, and when DBSCAN didn't work I tried other methods.

```
mins = 4
predictors = ["HP", "Attack", "Defense", "Speed", "Sp. Atk", "Sp. Def",
"Total"]
nn = NearestNeighbors(n_neighbors = mins + 1)
X = df.copy()
```

```
X = X[predictors]
z = StandardScaler()
X["predictors"] = z.fit(X[predictors])
nn.fit(X[predictors])
distances, neighbors = nn.kneighbors(X[predictors])
distances
distances = np.sort(distances[:, mins], axis = 0)
distances pokemon = pd.DataFrame({"distances": distances,
                             "index": list(range(0,len(distances)))})
plt = (ggplot(distances_pokemon, aes(x = "index", y = "distances")) +
geom_line(color = "white", size = 2) + theme_minimal() +
 labs(title = "Elbow Method for Choosing eps") +
 theme(panel grid minor = element blank(),
      rect = element rect(fill = "#202124ff"),
     axis_text = element_text(color = "white"),
      axis title = element text(color = "white"),
      plot title = element text(color = "white"),
      panel border = element line(color = "darkgray"),
      plot background = element rect(fill = "#202124ff")
      ))
ggsave(plot=plt, filename='elbow.png', dpi=300)
plt
```

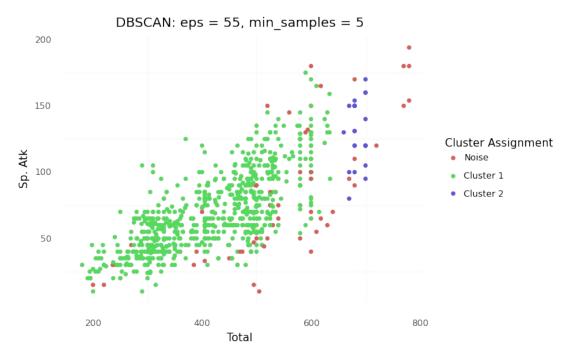


```
<ggplot: (8752395868925)>

df1 = DBSCAN(eps = 50, min_samples = 5).fit(X[predictors])
labsList = ["Noise"]
labsList = labsList + ["Cluster " + str(i) for i in
range(1,len(set(df1.labels_)))]

X["assignments"] = df1.labels_

(ggplot(X, aes(x = "Total", y = "Sp. Atk", color =
"factor(assignments)"))
+ geom_point()
+ theme_minimal()
+ scale_color_discrete(name = "Cluster Assignment", labels = labsList)
+ theme(panel_grid_major = element_blank())
+ labs(title = "DBSCAN: eps = 55, min_samples = 5"))
```



```
<ggplot: (8752400466557)>
ss_dbscan = X.loc[(X.assignments >= 0)]
print("SILHOUETTE:", silhouette_score(ss_dbscan[["Total", "Sp. Atk"]],
ss_dbscan["assignments"]))
SILHOUETTE: 0.4607780894566231
```

Here I used a Gaussian Mixture model to determine the probbility that each Pokemon is a Legendary, and I determined that it was ineffective because of how random the model was.

```
EM = GaussianMixture(n_components = 2)
X = df.copy()
X = X[predictors]
z = StandardScaler()

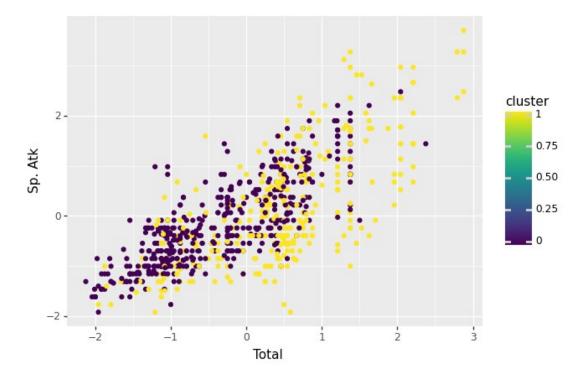
X[predictors] = z.fit_transform(X)
EM.fit(X)

cluster = EM.predict(X)

X["cluster"] = cluster

print((ggplot(X, aes(x = "Total", y = "Sp. Atk", color = "cluster")) +
geom_point()))

print("SILHOUETTE: ", silhouette_score(X, cluster))
```

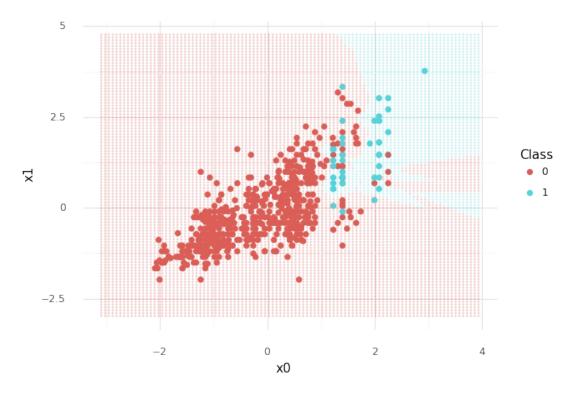


<ggplot: (8752396197761)>
SILHOUETTE: 0.17481438594257187

Here I used KNN and determined that it was super efficient because it tracked how many neighbors each Pokemon had.

```
predictors = ["Total", "Sp. Atk"]
X = df[predictors]
y = df["Legendary"]
n_neighbors = 5
knn = KNeighborsClassifier(n_neighbors = n_neighbors)
X_train, X_test, y_train, y_test = train_test_split(X,y, test_size = 0.2)
z = StandardScaler()
z.fit(X_train)
X_train[predictors] = z.transform(X_train)
X_test[predictors] = z.transform(X_test)
knn.fit(X_train,y_train)
X_train
import matplotlib.pyplot as plt
from matplotlib.colors import ListedColormap
```

```
def plotKNN2D(Xdf,v,k):
    # X can only have 2 dimensions because of plotting
    print("----")
    print(Xdf.columns)
    Xdf.columns = ["x0", "x1"]
    #grab the range of features for each feature
    x0_range = np.linspace(min(Xdf["x0"]) - np.std(Xdf["x0"]),
                           max(Xdf["x0"]) + np.std(Xdf["x0"]), num =
100)
    x1 range = np.linspace(min(Xdf["x1"]) - np.std(Xdf["x1"]),
                           max(Xdf["x1"]) + np.std(Xdf["x1"]), num =
100)
    #get all possible points on graph
    x0 = np.repeat(x0 range, 100)
    x1 = np.tile(x1 range, 100)
    x grid = pd.DataFrame(\{"x0": x0, "x1": x1\})
    #build model
    knn = KNeighborsClassifier(n neighbors = k)
    knn.fit(Xdf,y)
    # bredict all background points
    p = knn.predict(x grid)
    x grid["p"] = p #add to dataframe
    #build the plot
    bound = (ggplot(x grid, aes(x = "x0", y = "x1", color =
"factor(p)")) +
                 geom point(alpha = 0.2, size = 0.2) + theme minimal()
                 scale color discrete(name = "Class") +
                 geom point(data = Xdf, mapping = aes(x = "x0", y =
"x1", color = "factor(y)"), size = 2))
    print(bound)
plotKNN2D(X train, y train, k = n neighbors)
Index(['Total', 'Sp. Atk'], dtype='object')
```



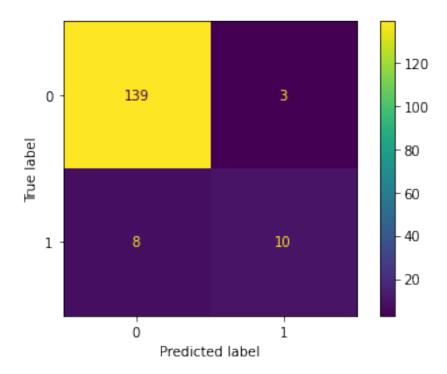
<ggplot: (8752403684845)>

print("ACCURACY:",knn.score(X_test,y_test))

ACCURACY: 0.93125

plot_confusion_matrix(knn, X_test, y_test)

<code><sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay</code> at 0x7f5d3b99bd90>



3)

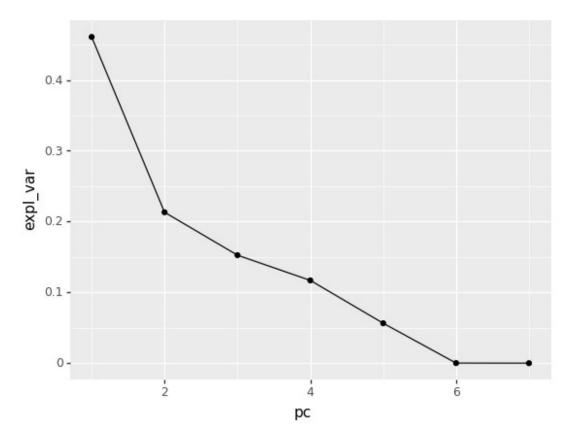
When using HP, Attack, Defense, Speed, Special attack, Special Defense, and Total as my predictors to determine what type a Pokemon is, can I use PCA to create a more efficient model?

In this model I determined that when using PCA to remove 4 of my predictors, and using a regular Logistic Regression both models produced very inaccurate predictions.

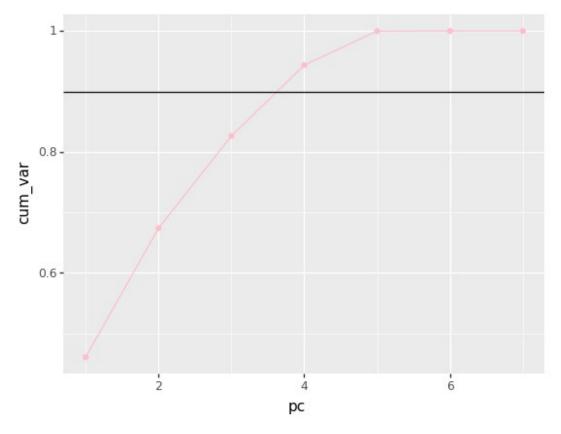
In my PCA model I determined that I can keep over 90% accuracy by keeping only 4 predictors and prodiced a model with an R2 score of -1 for train and -1.47 for test and an MSE of 59 for train and 71 for test, showing that the varience of the data could not be explained by the model, and that the errors squared were, and that the regression line was nowere near my points.

In my Logistic Regression model I was able to determine that the R2 score was -0.89 for train and -1.4 for test and the MSE was 56 for train and 69.67 for test, so this was also very inaccurate. When graphing my coefficients I determined that none of the coefficients were above an absolite value of 0.1, meaning that none of the predictors were impactful at all.

```
df["Type 1"], test size=0.2)
z = StandardScaler()
X_train[predictors] = z.fit_transform(X_train[predictors])
X test[predictors] = z.transform(X test[predictors])
lr = LogisticRegression()
lr.fit(X train, y train)
pca = PCA()
pca.fit(X_train[features], y_train)
pcaDF = pd.DataFrame({"expl var" :
                      pca.explained_variance_ratio_,
                      "pc": range(1,8),
                      "cum var":
                      pca.explained_variance_ratio_.cumsum()
pcaDF.head()
   expl var
            рс
                cum var
0 0.482405
            1 0.482405
1 0.197129
             2 0.679534
2 0.145420
            3 0.824954
3 0.117848
            4 0.942801
4 0.057047
             5 0.999848
(ggplot(pcaDF, aes(x = "pc", y = "expl_var")) + geom_line() +
geom_point())
```



```
<ggplot: (8752400579529)>
(ggplot(pcaDF, aes(x = "pc", y = "cum_var")) + geom_line(color = "pink") +
geom_point(color = "pink") + geom_hline(yintercept = 0.90))
```



<ggplot: (8752400564509)>

Here I used PCA to create my model with only 4 of the predictors.

```
pcomps4Train = pca.transform(X_train)
pcomps4Test = pca.transform(X_test)
pcomps4Train = pd.DataFrame(pcomps4Train[:, 0:4])
pcomps4Test = pd.DataFrame(pcomps4Test[:, 0:4])
lr1 = LogisticRegression()
lr1.fit(pcomps4Train, y_train)

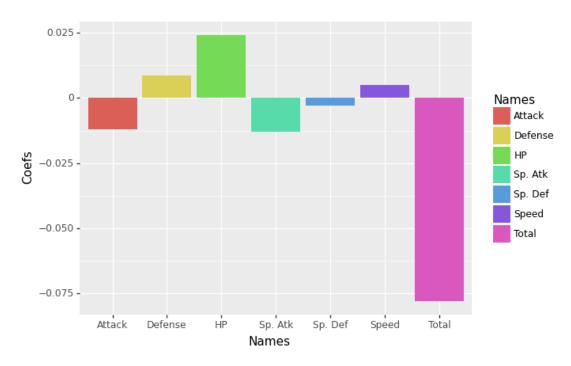
print("MSE Train:", mean_squared_error(y_train,
lr1.predict(pcomps4Train)))
print("MSE Test:", mean_squared_error(y_test,
lr1.predict(pcomps4Test)))

print("R2 Train:",r2_score(y_train, lr1.predict(pcomps4Train)))
print("R2 Test:",r2_score(y_test, lr1.predict(pcomps4Test)))

MSE Train: 59.4140625
MSE Test: 71.28125
R2 Train: -1.0009078741266477
```

R2 Train: -1.0009078741266477 R2 Test: -1.4690957112857492 Here I used all my predictors for a Logistic Regression.

```
yPredTrain = lr.predict(X train)
yPredTest = lr.predict(X test)
print("MSE Train:" , mean_squared_error(y_train,yPredTrain ))
print("MSE Test:" , mean_squared_error(y_test,yPredTest ))
print("R2 Train:",r2_score(y_train,yPredTrain ))
print("R2 Test:",r2_score(y_test,yPredTest ))
MSE Train: 56.1140625
MSE Test: 69.66875
R2 Train: -0.8897726359897515
R2 Test: -1.413240674590289
coef = pd.DataFrame({"Coefs": lr.coef_[0],
                       "Names": features})
coef
      Coefs
                Names
0 0.024047
                   HP
1 -0.011997
             Attack
2 0.008517 Defense
3 0.004989
                Speed
4 -0.013052 Sp. Atk
5 -0.002963 Sp. Def
                Total
6 -0.078013
(ggplot(coef, aes(x = "Names", y = "Coefs", fill = "Names")) +
geom bar(stat = "identity"))
```



<ggplot: (8752400496581)>