# Machine Learning Model Prediction

#### **Data Preparation**

Required Specific Libraries

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as matplt
from matplotlib import rcParams
import seaborn as sns
%matplotlib inline
import warnings
warnings.filterwarnings("ignore")
from \ sklearn.preprocessing \ import \ Standard Scaler, \ Label Encoder, \ Ordinal Encoder, \ One Hot Encoder
from sklearn.compose import make_column_transformer
from imblearn.over_sampling import SMOTE
from sklearn.model_selection import train_test_split
from \ sklearn. metrics \ import \ classification\_report, ConfusionMatrix Display, \ accuracy\_score, \ confusion\_matrix, \ precision\_score, \ recall\_score
from google.colab import drive
drive.mount('/content/drive')
 Trive already mounted at /content/drive; to attempt to forcibly remount, call drive.mount("/content/drive", force_remount=True).

    Import dataset

dataframe = pd.read_csv('/content/drive/My Drive/Colab Notebooks/Exam/LA4PSchools.csv')
dataframe['Year3_Writing_At_Risk'] = dataframe['Year3_Writing_At_Risk'].map({False: 0, True: 1})
Preprocess the data
```

dataframe.drop(["StudentID"],axis=1,inplace=True)

### For Encoding Categorical data

```
change_values = {
  'Gender': {
     'Male': 0,
     'Female': 1
}
dataframe.replace(change_values, inplace=True)
dataframe.head()
```

₹		Gender	Year_02	Kinder_Age	Disability	NCCD- Funded	01.SES	02.SES	NumSibling	SiblingOrder	NumAbvYear9	•••	HRSIW- 01-SOY	Counting- 01	Coun
	0	0	2020	5.5	Disability_Non- disable	0	104	104	3	3	2		49	4	
	1	1	2018	5.8	Disability_Non- disable	0	112	112	2	2	2		37	2	
	2	0	2021	5.9	Disability_Non- disable	0	120	109	2	2	2		30	2	
	3	0	2021	5.7	Disability_Non- disable	0	95	93	2	1	0		30	2	
	4	0	2021	5.8	Disability_Non- disable	0	98	98	1	1	2		32	2	
	5 ro	ws × 33 c	columns												
	4														•
dataframe_copy = dataframe															
disability_transformer = make_column_transformer(															

```
disability_transformer = make_column_transformer(
   (OneHotEncoder(), ['Disability']),
   remainder='passthrough',
   verbose_feature_names_out=False
)

dataframe = dataframe_copy

work_type_transformed = disability_transformer.fit_transform(dataframe)
dataframe = pd.DataFrame(work_type_transformed, columns=disability_transformer.get_feature_names_out())
dataframe.head(5)
```

Disability_	_Disability_Cognitive	Disability_Disability_Non- disable	Disability_Disability_Physical	Disability_Disability_Sensory	Disabili1
0	0.0	1.0	0.0	0.0	
1	0.0	1.0	0.0	0.0	
2	0.0	1.0	0.0	0.0	
3	0.0	1.0	0.0	0.0	
4	0.0	1.0	0.0	0.0	
5 rows × 37 colum	nns				
4					<b>&gt;</b>

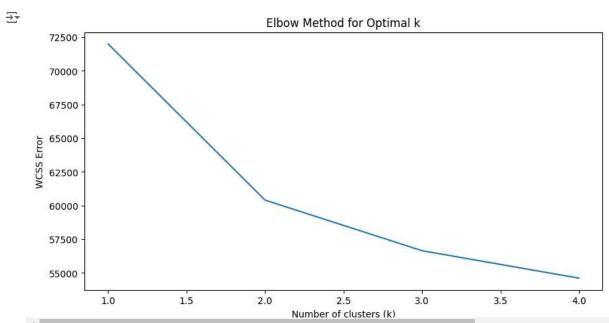
# Model Training

₹

```
depent_axis = dataframe["Year3_Writing_At_Risk"]
independent_axis = dataframe.drop(["Year3_Writing_At_Risk"],axis=1)
```

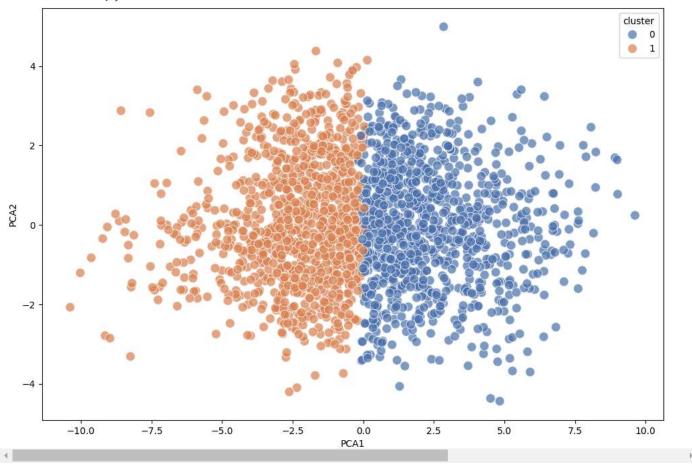
# Scaling Dataset from Standardize

```
-0.09012398, 0.26241365],
[ 1.80676294, -1.49365919, -0.18617505, ..., -0.22284613,
              1.10357108, 2.85415341]])
from sklearn.cluster import KMeans
import matplotlib.pyplot as plt
# Determine the number of clusters (k) using the Elbow Method
wcss = [] # Within-cluster sum of squares
for i in range(1, 5):
    kmeans = KMeans(n_clusters=i, random_state=0)
    kmeans.fit(independent_axis) # Using the features
    wcss.append(kmeans.inertia_)
# Plotting the elbow curve
plt.figure(figsize=(10, 5))
plt.plot(range(1, 5), wcss)
plt.title('Elbow Method for Optimal k')
plt.xlabel('Number of clusters (k)')
plt.ylabel('WCSS Error')
plt.show()
```

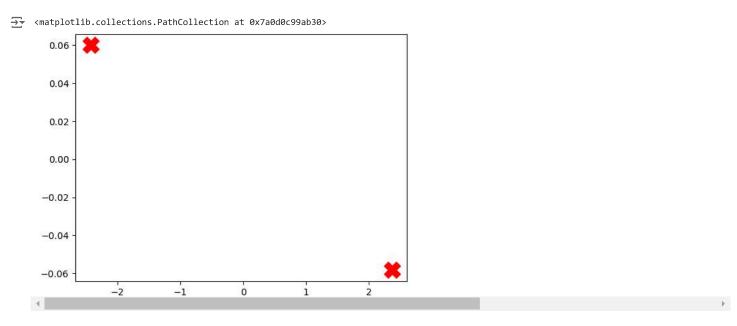


```
WCSS
```

```
[71999.999999997, 60408.88619173092, 56653.825553083385, 54615.53781676896]
# Assuming optimal k is chosen, say k=2
optimal_k = 2
kmeans = KMeans(n_clusters=optimal_k, random_state=42)
# Fit and predict clusters
clusters = kmeans.fit_predict(independent_axis)
dataframe['cluster'] = clusters
# Optional: Use PCA to reduce dimensions to 2 for visualization
from sklearn.decomposition import PCA
pca = PCA(n_components=2)
reduced_data = pca.fit_transform(independent_axis)
visualization_df = pd.DataFrame(reduced_data, columns=['PCA1', 'PCA2'])
visualization_df['cluster'] = clusters
# Plotting clusters
plt.figure(figsize=(12, 8))
sns.scatterplot(data=visualization_df, x='PCA1', y='PCA2', hue='cluster', palette='deep', s=100, alpha=0.7)
```



# Plotting centroids
centroids = pca.transform(kmeans.cluster\_centers\_) # Transform centroids to PCA space
plt.scatter(centroids[:, 0], centroids[:, 1], s=300, c='red', marker='X', label='Centroids')



```
labels = kmeans.labels_
correct_labels = sum(depent_axis == labels)

print("Result: %d out of %d samples were correctly labeled." % (correct_labels, depent_axis.size))

Result: 1373 out of 2000 samples were correctly labeled.
```

 $\verb|print('Accuracy score: \{0:0.2f\}'. format(correct\_labels/float(depent\_axis.size)))| \\$ 

→ Accuracy score: 0.69