



Assessment Report
on
“Health Risk Classification”
submitted as partial fulfillment for the award of
BACHELOR OF TECHNOLOGY
DEGREE

SESSION 2024-25

in
CSE(AI)

By

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Introduction :

In today's world, predicting health risk is important to take precautions early. This project focuses on predicting whether a person is at low, medium, or high health risk by analyzing their Body Mass Index (BMI), exercise hours per week, and junk food consumption frequency. We used machine learning to build a model that can classify these risk levels automatically.

Methodology :

1. Dataset: We used a dataset containing health information like BMI, exercise hours, junk food frequency, and risk level.
2. Preprocessing: We converted the risk level (Low/Medium/High) into numeric values.
3. Model Used: We used the Random Forest Classifier from sklearn.
4. Train/Test Split: 80% of the data was used for training and 20% for testing.
5. Evaluation: We used metrics like accuracy, precision, recall, and a confusion matrix heatmap to check model performance.
6. Clustering: We also applied KMeans clustering to group people into health risk clusters without using labels.

Code :

```
# === STEP 1: Import Required Libraries ===  
  
import pandas as pd  
import matplotlib.pyplot as plt  
import seaborn as sns  
  
from sklearn.model_selection import train_test_split  
from sklearn.ensemble import RandomForestClassifier  
from sklearn.metrics import confusion_matrix, accuracy_score,  
precision_score, recall_score, classification_report  
  
from sklearn.cluster import KMeans  
from sklearn.preprocessing import StandardScaler  
from sklearn.metrics import silhouette_score  
  
# === STEP 2: Upload and Load Dataset ===  
  
from google.colab import files  
uploaded = files.upload()  
  
# Automatically pick the uploaded file  
file_name = next(iter(uploaded))  
df = pd.read_csv(file_name)
```

```
# === STEP 3: Show First Few Rows ===
```

```
print("🔍 Sample of Data:")
```

```
print(df.head())
```

```
# === STEP 4: Classification (Supervised Learning) ===
```

```
print("\n🧠 Classification: Risk Level Prediction")
```

```
# Convert risk_level to numerical: low=0, medium=1, high=2
```

```
df['risk_level'] = df['risk_level'].map({'low': 0, 'medium': 1, 'high': 2})
```

```
# Features and target
```

```
X = df[['bmi', 'exercise_hours', 'junk_food_freq']]
```

```
y = df['risk_level']
```

```
# Train-test split
```

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2,  
random_state=42)
```

```
# Train classifier
```

```
clf = RandomForestClassifier(random_state=42)
```

```
clf.fit(X_train, y_train)
```

```
# Predict
```

```
y_pred = clf.predict(X_test)
```

```
# Evaluation Metrics
```

```
accuracy = accuracy_score(y_test, y_pred)
```

```
precision = precision_score(y_test, y_pred, average='macro',  
zero_division=0)
```

```
recall = recall_score(y_test, y_pred, average='macro', zero_division=0)
```

```
print("\n 📊 Evaluation Metrics:")
```

```
print(f"Accuracy : {accuracy:.2f}")
```

```
print(f"Precision: {precision:.2f}")
```

```
print(f"Recall : {recall:.2f}")
```

```
print("\nDetailed Report:")
```

```
print(classification_report(y_test, y_pred, target_names=["Low", "Medium",  
"High"]))
```

```
# Confusion Matrix Heatmap
```

```
cm = confusion_matrix(y_test, y_pred)
```

```
plt.figure(figsize=(8, 6))
```

```
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues',
```

```
            xticklabels=["Low", "Medium", "High"],
```

```
            yticklabels=["Low", "Medium", "High"])
```

```
plt.title("Confusion Matrix Heatmap")
plt.xlabel("Predicted")
plt.ylabel("Actual")
plt.show()
```

```
# === STEP 5: Clustering (Unsupervised Learning) ===
```

```
print("\n🔍 Clustering (KMeans) for Health Profile Segmentation")
```

```
# Remove target column for clustering
```

```
X_cluster = df[['bmi', 'exercise_hours', 'junk_food_freq']]
```

```
# Scale the features
```

```
scaler = StandardScaler()
```

```
X_scaled = scaler.fit_transform(X_cluster)
```

```
# Apply KMeans clustering (let's try 3 clusters)
```

```
kmeans = KMeans(n_clusters=3, random_state=42, n_init=10)
```

```
clusters = kmeans.fit_predict(X_scaled)
```

```
# Add cluster labels to dataframe
```

```
df['cluster'] = clusters
```

```
# Evaluate clustering
```


```
sil_score = silhouette_score(X_scaled, clusters)
print(f"\nSilhouette Score (Cluster Quality): {sil_score:.2f}")

# Visualize clusters
plt.figure(figsize=(8, 6))
sns.scatterplot(x=df['bmi'], y=df['exercise_hours'], hue=clusters,
palette='Set1')
plt.title("KMeans Clustering: Health Segmentation")
plt.xlabel("BMI")
plt.ylabel("Exercise Hours/Week")
plt.legend(title="Cluster")
plt.show()
```



Screenshot of Output :


- **health_risk (1).csv**(text/csv) - 2881 bytes, last modified: 4/22/2025 - 100% done

Saving health_risk (1).csv to health_risk (1) (1).csv

 Sample of Data:

	bmi	exercise_hours	junk_food_freq	risk_level
0	28.730279	13	1	high
1	31.301442	12	4	medium
2	32.549043	9	0	medium
3	30.463670	2	1	medium
4	28.431755	2	1	low

 Classification: Risk Level Prediction

 Evaluation Metrics:

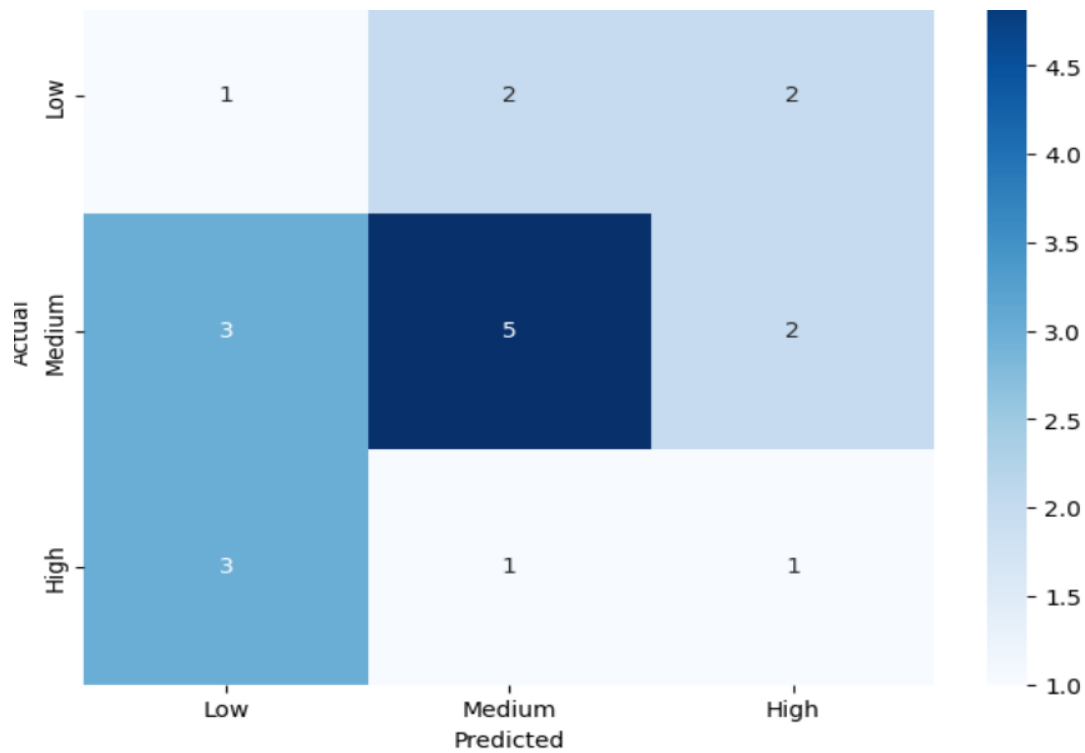
Accuracy : 0.35

Precision: 0.32

Recall : 0.30

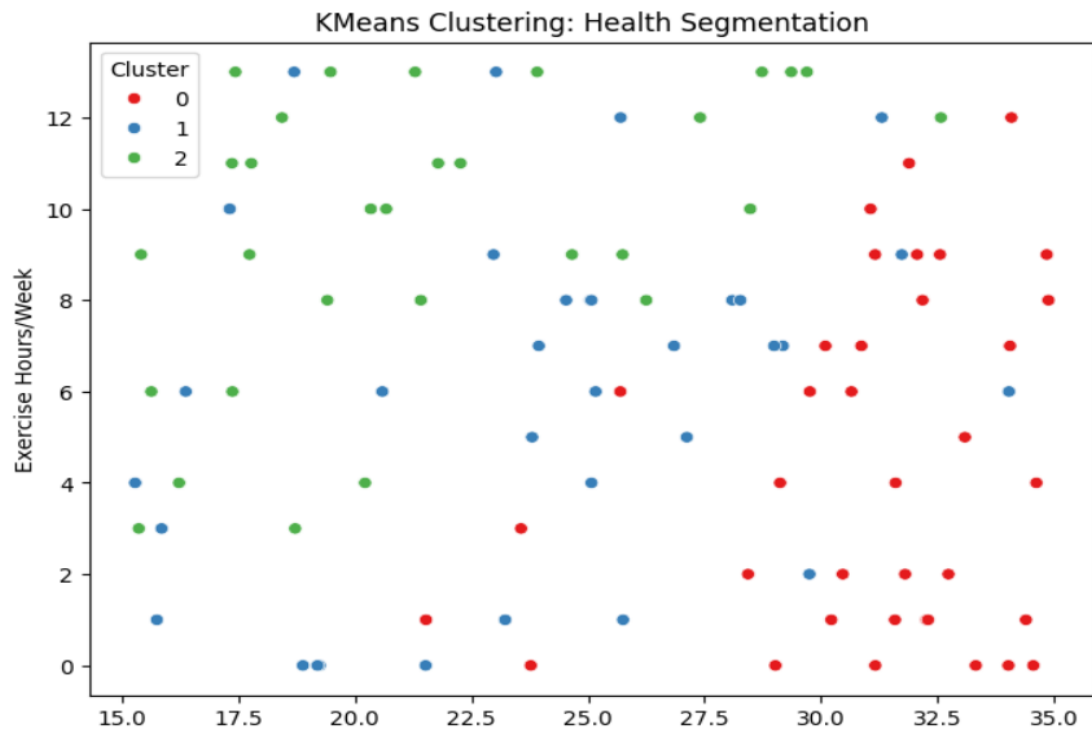
Detailed Report:

	precision	recall	f1-score	support
Low	0.14	0.20	0.17	5
Medium	0.62	0.50	0.56	10
High	0.20	0.20	0.20	5
accuracy			0.35	20
macro avg	0.32	0.30	0.31	20
weighted avg	0.40	0.35	0.37	20



🔍 Clustering (KMeans) for Health Profile Segmentation

Silhouette Score (Cluster Quality): 0.28



References/Credits :

- . Dataset: Provided by faculty / or custom-created
- . Libraries: scikit-learn, pandas, matplotlib, seaborn
- . Colab: Google Colab used for cloud-based execution
- . Guide: Assistance provided by AI tools and online documentation

