numerical_data.head()

```
import pandas as pd
import numpy as np
import scipy.stats as stats
import statsmodels.api as sm
import seaborn as sns
import matplotlib.pyplot as plt

data = pd.read_csv('./Q 7- Copy.csv')

# View the first few rows of the dataset
# print(data.head())
data.head()

# Select numerical columns for analysis
# numerical_data = data.select_dtypes(include=np.number)
```

∓ Quality Physical BMI Sleep Person Sleep Stress Blood Heart Daily Gender Age Occupation of **Activity** Duration Steps Disorder TD Level Category Pressure Rate Sleep Level Software 0 27 4200 1 Male 6.1 6 42 6 Overweight 126/83 77 NaN Engineer 1 2 Male 28 Doctor 6.2 6 60 8 Normal 125/80 10000 NaN 75 2 3 10000 Male 28 Doctor 6.2 60 Normal 125/80 75 NaN Sales Sleep 3 5.9 30 8 Obese 140/90 3000 4 Male 28 85 Representative Apnea Sales Sleep 4 5 Male 5.9 4 30 8 Obese 140/90 85 3000

```
# Importing necessary libraries
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.preprocessing import StandardScaler, LabelEncoder
from sklearn.decomposition import PCA
# Load the dataset (assuming the dataset is in CSV format)
# Replace 'dataset.csv' with the actual file path
df = pd.read_csv('./Q 7- Copy.csv')
# Display the first few rows of the dataset
print(df.head())
# Data Preprocessing
# Handle missing values by filling with median or mean (for simplicity, using mean here)
# df.fillna(df.mean(), inplace=True)
# Encode categorical variables
label_encoders = {}
for column in ['Gender', 'Sleep Duration']:
    le = LabelEncoder()
    df[column] = le.fit_transform(df[column])
    label_encoders[column] = le
# Standardize numerical columns
numerical_features = ['Age', 'Sleep Duration', 'Physical Activity Level', 'Stress Level', 'Quality of Sleep']
scaler = StandardScaler()
df[numerical_features] = scaler.fit_transform(df[numerical_features])
# Apply PCA
pca = PCA(n_components=2) # We reduce to 2 components for visualization purposes
```

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pca_components = pca.fit_transform(df[numerical_features])
# Create a DataFrame with PCA components
pca_df = pd.DataFrame(data=pca_components, columns=['PC1', 'PC2'])
# Visualize the explained variance ratio
plt.figure(figsize=(8, 5))
plt.plot(range(1, len(pca.explained_variance_ratio_) + 1), pca.explained_variance_ratio_, marker='o', linestyle='--')
plt.title('Explained Variance Ratio')
plt.xlabel('Principal Components')
plt.ylabel('Variance Explained')
plt.show()
# Scatter plot of the PCA results
plt.figure(figsize=(10, 7))
sns.scatterplot(x='PC1', y='PC2', data=pca_df, hue=df['Gender'], palette='viridis', alpha=0.6)
plt.title('PCA of Sleep Health Factors')
plt.xlabel('Principal Component 1')
plt.ylabel('Principal Component 2')
plt.show()
# Display PCA components (loading scores)
components df = pd.DataFrame(data=pca.components , columns=numerical features, index=['PC1', 'PC2'])
print("\nPCA Components:")
print(components df)
# Interpretation of findings
# The PCA components can be interpreted by examining the loading scores to see which features contribute most to each components
# For example, higher values in 'PC1' might indicate the importance of sleep duration and stress level.
# Higher values in 'PC2' could highlight physical activity and caffeine intake.
# Additional: Saving the PCA results for further analysis
pca_df['Gender'] = df['Gender']
# pca_df.to_csv('pca_results.csv', index=False)
```





