

```
#importing libraries
```

```
import numpy as np
```

```
import pandas as pd
```

```
import matplotlib.pyplot as plt
```

```
import seaborn as sns
```

```
#loading the dataset
```

```
df = pd.read_csv('Sleep_health_and_lifestyle_dataset.csv')
```

```
df.head()
```

| | Person ID | Gender | Age | Occupation | Sleep Duration \ |
|---|-----------|--------|-----|----------------------|------------------|
| 0 | 1 | Male | 27 | Software Engineer | 6.1 |
| 1 | 2 | Male | 28 | Doctor | 6.2 |
| 2 | 3 | Male | 28 | Doctor | 6.2 |
| 3 | 4 | Male | 28 | Sales Representative | 5.9 |
| 4 | 5 | Male | 28 | Sales Representative | 5.9 |

| | Quality of Sleep Category \ | Physical Activity Level | Stress Level | BMI |
|------------|-----------------------------|-------------------------|--------------|-----|
| 0 | | 6 | 42 | 6 |
| Overweight | | | | |
| 1 | | 6 | 60 | 8 |
| Normal | | | | |
| 2 | | 6 | 60 | 8 |
| Normal | | | | |
| 3 | | 4 | 30 | 8 |
| Obese | | | | |
| 4 | | 4 | 30 | 8 |
| Obese | | | | |

| | Blood Pressure | Heart Rate | Daily Steps | Sleep Disorder |
|---|----------------|------------|-------------|----------------|
| 0 | 126/83 | 77 | 4200 | NaN |
| 1 | 125/80 | 75 | 10000 | NaN |
| 2 | 125/80 | 75 | 10000 | NaN |
| 3 | 140/90 | 85 | 3000 | Sleep Apnea |
| 4 | 140/90 | 85 | 3000 | Sleep Apnea |

```
#checking for missing values
```

```
df.isnull().sum()
```

| | |
|-------------------------|---|
| Person ID | 0 |
| Gender | 0 |
| Age | 0 |
| Occupation | 0 |
| Sleep Duration | 0 |
| Quality of Sleep | 0 |
| Physical Activity Level | 0 |
| Stress Level | 0 |
| BMI Category | 0 |
| Blood Pressure | 0 |

```
Heart Rate          0
Daily Steps         0
Sleep Disorder      219
dtype: int64
```

```
#replacing the null values with 'None' in the column 'Sleep Disorder'
df['Sleep Disorder'].fillna('None', inplace=True)
```

```
#drop column Person ID
df.drop('Person ID', axis=1, inplace=True)
```

```
# print the number of unique values for each column to get a quick
sense of data variety
print("Unique values in each column:")
for col in df.columns:
    unique_count = df[col].nunique() # count unique entries in the
column
    print(f"{col}: {unique_count}") # show column name and its
unique count
```

Unique values in each column:

```
Gender: 2
Age: 31
Occupation: 11
Sleep Duration: 27
Quality of Sleep: 6
Physical Activity Level: 16
Stress Level: 6
BMI Category: 4
Heart Rate: 19
Daily Steps: 20
Sleep Disorder: 3
systolic_bp: 18
diastolic_bp: 17
```

```
# clean any sneaky spaces in column names
df.columns = df.columns.str.strip()
```

```
# split bp if it's there, else warn me
if 'Blood Pressure' in df.columns:
    df[['systolic_bp', 'diastolic_bp']] = df['Blood
Pressure'].str.split('/', expand=True)
    df.drop('Blood Pressure', axis=1, inplace=True)
else:
    print("no 'Blood Pressure' column found")
```

```
no 'Blood Pressure' column found
```

```
df['BMI Category'] = df['BMI Category'].replace('Normal Weight',
'Normal')
```

```
df.head()
```

| | Gender | Age | Occupation | Sleep Duration | Quality of Sleep \ |
|---|--------|-----|------------|----------------|--------------------|
| 0 | 1 | 27 | 9 | 6.1 | 6 |
| 1 | 1 | 28 | 1 | 6.2 | 6 |
| 2 | 1 | 28 | 1 | 6.2 | 6 |
| 3 | 1 | 28 | 6 | 5.9 | 4 |
| 4 | 1 | 28 | 6 | 5.9 | 4 |

| | Physical Activity Level | Stress Level | BMI Category | Blood Pressure |
|---|-------------------------|--------------|--------------|----------------|
| 0 | 42 | 6 | 3 | 126/83 |
| 1 | 60 | 8 | 0 | 125/80 |
| 2 | 60 | 8 | 0 | 125/80 |
| 3 | 30 | 8 | 2 | 140/90 |
| 4 | 30 | 8 | 2 | 140/90 |

| | Heart Rate | Daily Steps | Sleep Disorder |
|---|------------|-------------|----------------|
| 0 | 77 | 4200 | 1 |
| 1 | 75 | 10000 | 1 |
| 2 | 75 | 10000 | 1 |
| 3 | 85 | 3000 | 2 |
| 4 | 85 | 3000 | 2 |

```
#unique values from categorical columns
```

```
print(df.Occupation.unique())
```

```
print('\n')
```

```
print(df['BMI Category'].unique())
```

```
print('\n')
```

```
print(df['Sleep Disorder'].unique())
```

```
['Software Engineer' 'Doctor' 'Sales Representative' 'Teacher' 'Nurse'  
'Engineer' 'Accountant' 'Scientist' 'Lawyer' 'Salesperson' 'Manager']
```

```
['Overweight' 'Normal' 'Obese' 'Normal Weight']
```

```
['None' 'Sleep Apnea' 'Insomnia']
```

```
fig, ax = plt.subplots(3,3, figsize=(20,10))
```

```
sns.countplot(x='Gender', data=df, ax=ax[0,0], hue='Gender',  
palette='pastel', legend=False)
```

```
sns.histplot(x='Age', data=df, ax=ax[0,1], bins=10, color='#f7cac9')
```

```
# soft pink coral-ish
```

```

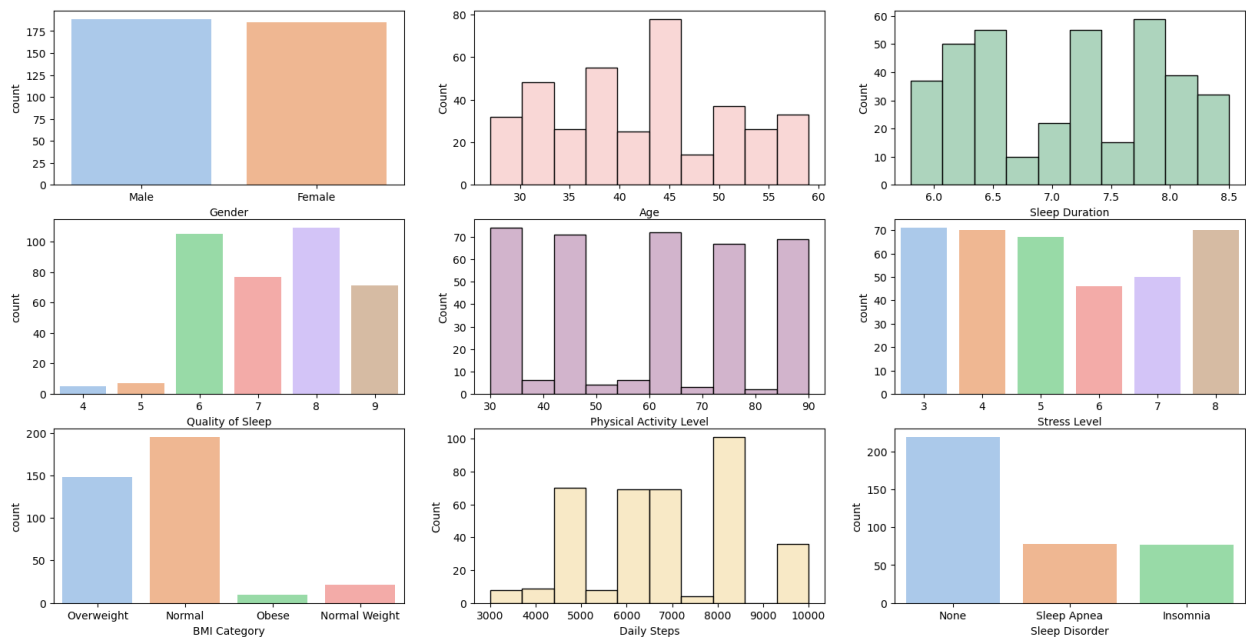
sns.histplot(x='Sleep Duration', data=df, ax=ax[0,2], bins=10,
color='#92c6a7') # soft green

sns.countplot(x='Quality of Sleep', data=df, ax=ax[1,0], hue='Quality
of Sleep', palette='pastel', legend=False)
sns.histplot(x='Physical Activity Level', data=df, ax=ax[1,1],
bins=10, color='#c49bbb') # muted orchid-ish
sns.countplot(x='Stress Level', data=df, ax=ax[1,2], hue='Stress
Level', palette='pastel', legend=False)

sns.countplot(x='BMI Category', data=df, ax=ax[2,0], hue='BMI
Category', palette='pastel', legend=False)
sns.histplot(x='Daily Steps', data=df, ax=ax[2,1], bins=10,
color='#f6e2b3') # light gold
sns.countplot(x='Sleep Disorder', data=df, ax=ax[2,2], hue='Sleep
Disorder', palette='pastel', legend=False)

<Axes: xlabel='Sleep Disorder', ylabel='count'>

```



```

pink_pastel = ['#f8bbd0', '#f48fb1', '#f06292'] # 3 pastel pink
shades

```

```

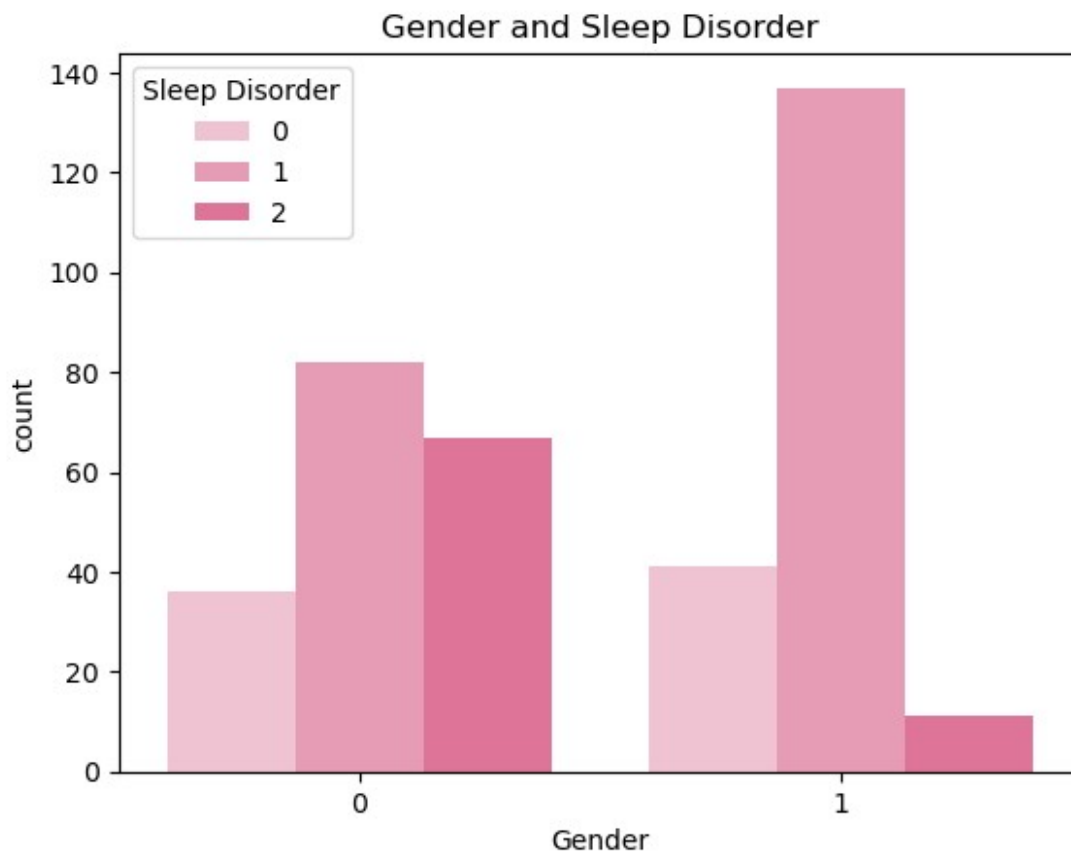
sns.countplot(x='Gender', data=df, hue='Sleep Disorder',
palette=pink_pastel).set_title('Gender and Sleep Disorder')

```

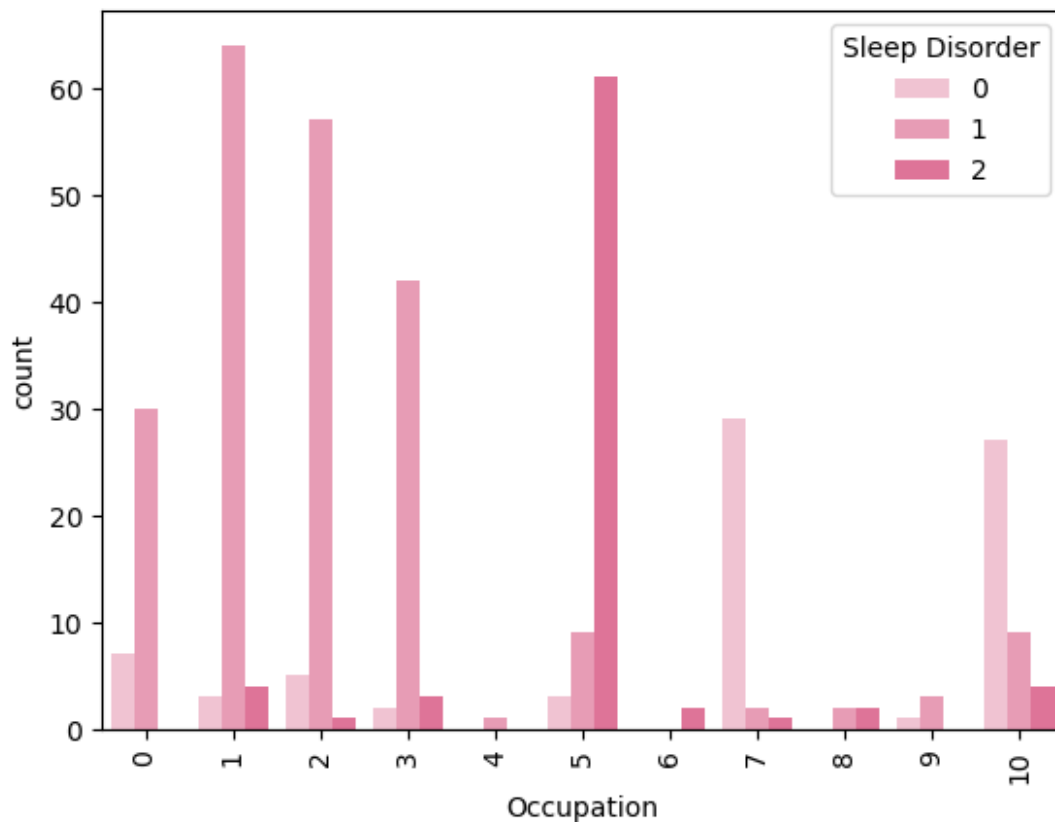
```

Text(0.5, 1.0, 'Gender and Sleep Disorder')

```



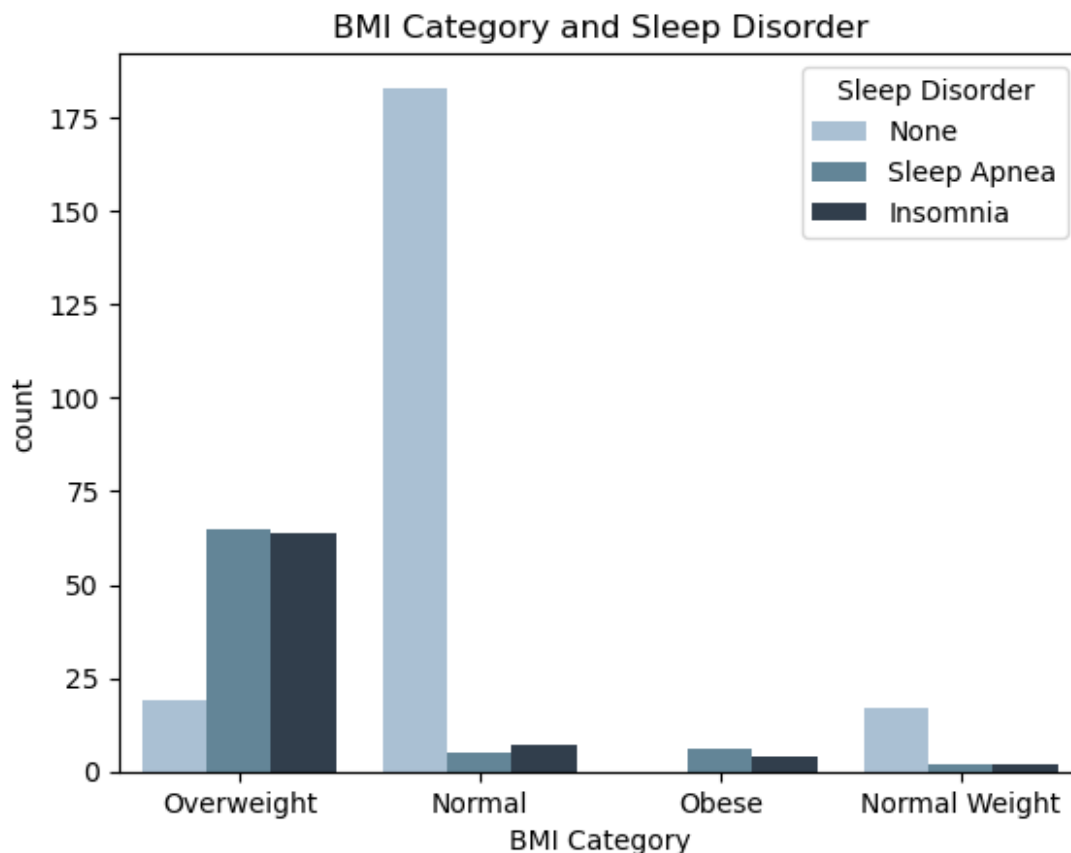
```
pink_pastel = ['#f8bbd0', '#f48fb1', '#f06292'] # 3 soft pink shades
ax = sns.countplot(x='Occupation', data=df, hue='Sleep Disorder',
palette=pink_pastel)
plt.xticks(rotation=90)
plt.legend(title='Sleep Disorder')
<matplotlib.legend.Legend at 0x16a268200>
```



```
blue_shades = ['#a3c1da', '#5a87a0', '#2c3e50'] # light to dark blue shades
```

```
sns.countplot(x='BMI Category', hue='Sleep Disorder', data=df,
palette=blue_shades).set_title('BMI Category and Sleep Disorder')
```

```
Text(0.5, 1.0, 'BMI Category and Sleep Disorder')
```



```
from sklearn.preprocessing import LabelEncoder

vars = ['Gender', 'Occupation', 'BMI Category', 'Sleep Disorder']
for i in vars:
    label_encoder = LabelEncoder() # make a fresh encoder each loop
    df[i] = label_encoder.fit_transform(df[i])
    print(i, ': ', df[i].unique())

Gender : [1 0]
Occupation : [ 9  1  6 10  5  2  0  8  3  7  4]
BMI Category : [3 0 2 1]
Sleep Disorder : [1 2 0]

plt.figure(figsize=(20,16))
sns.heatmap(df.select_dtypes(include='number').corr(), annot=True,
            cmap='vlag')

<Axes: >
```

