Final Project

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```
library(tidyverse)
library(broom)
library(tidyr)
library(dplyr)
library(modelr)
library(boot)
library(tidyr)
library(ggplot2)
library(ggmosaic)
library(dplyr)
library(readr)
library(class)
library(caret)
library(infer)
```

#Intro

```
library(tidyr)
library(ggplot2)
library(ggmosaic)
library(dplyr)
Sleep_health_and_lifestyle_dataset <- read_csv("Sleep_health_and_lifestyle_dataset.csv")
```

```
Sleep_health_and_lifestyle_dataset_renamed <- Sleep_health_and_lifestyle_dataset %>%
  rename( Duration = 'Sleep Duration',
          Stress = 'Stress Level',
          Physical = 'Physical Activity Level' ,
          Quality = 'Quality of Sleep' ,
          BMI= 'BMI Category' ,
          BPressure = 'Blood Pressure' ,
          HRate = 'Heart Rate' ,
          DSteps = 'Daily Steps' ,
          Disorder = 'Sleep Disorder' )
```

#Predictive Analysis

###Load the dataset

```
Sleep_health_and_lifestyle_dataset <- read_csv(file = "Sleep_health_and_lifestyle_dataset.csv"
  col_types = cols(
    'Person ID' = col_character(),
    'Age' = col_double(),
    'Sleep Duration' = col_double(),
    'Stress Level' = col_double(),
    'Physical Activity Level' = col_double(),
    'Quality of Sleep' = col_double(),
    'BMI Category' = col_character(),
    'Blood Pressure' = col_character(),
    'Heart Rate' = col_double(),
    'Daily Steps' = col_double(),
    'Sleep Disorder' = col_character()
  ))
```

Rename

```
Sleep_health_and_lifestyle_dataset_renamed <- Sleep_health_and_lifestyle_dataset %>%
  rename(ID = 'Person ID',
         Duration = 'Sleep Duration',
         Stress = 'Stress Level',
         Physical = 'Physical Activity Level',
         Quality = 'Quality of Sleep',
         BMI = 'BMI Category',
         BPressure = 'Blood Pressure',
         HRate = 'Heart Rate',
         DSteps = 'Daily Steps',
         Disorder = 'Sleep Disorder')
```

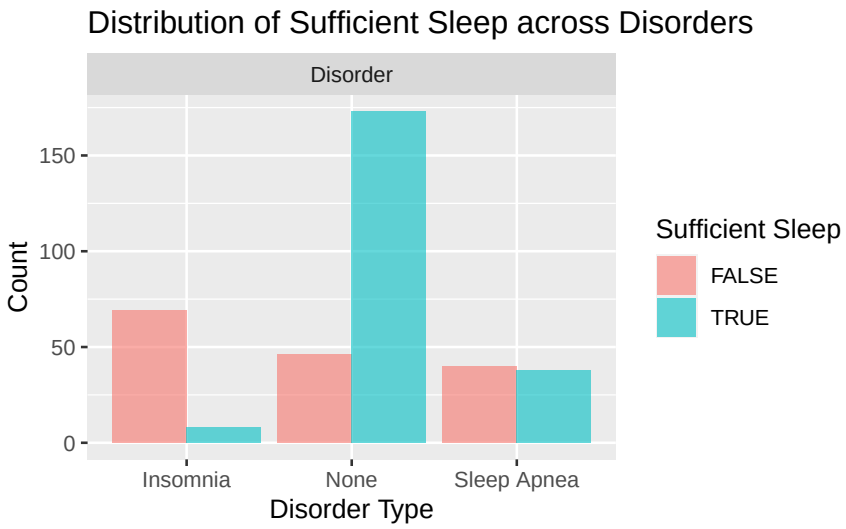
Parse Sleep Data

```
sleep_data <- Sleep_health_and_lifestyle_dataset_renamed %>%
    mutate(sufficient_sleep = as.logical(Duration >= 7.0))
```

Sleep Data Disorders

```
sleep_data %>%
  pivot_longer(cols = c(Disorder), names_to = "variable", values_to = "value") %>%
```

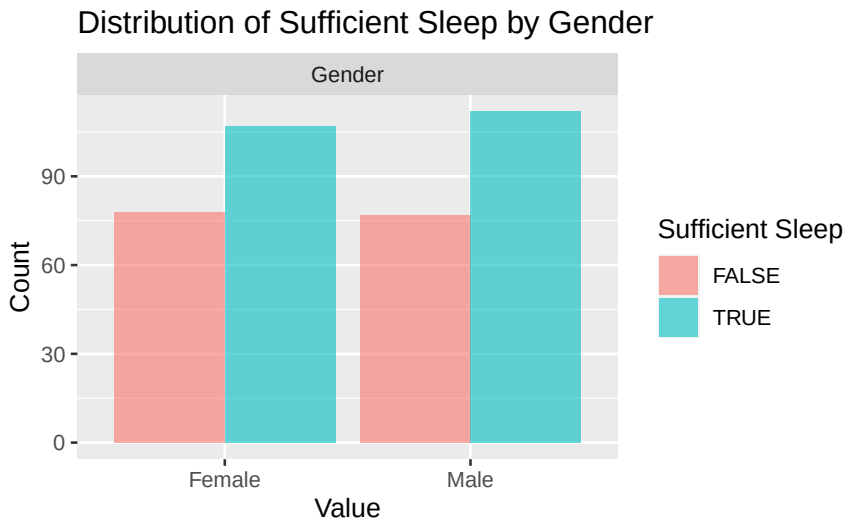
```
  group_by(variable, value, sufficient_sleep) %>%
  summarise(count = n()) %>%
  ggplot() +
  geom_bar(
    mapping = aes(x = value, y = count, fill = sufficient_sleep),
    position = "dodge",
    alpha = 0.6,
    stat = "identity"
  ) +
  facet_wrap(~ variable, scales = "free") +
  labs(title = "Distribution of Sufficient Sleep across Disorders",
       x = "Disorder Type",
       y = "Count",
       fill = "Sufficient Sleep")
```



Sleep Data Gender

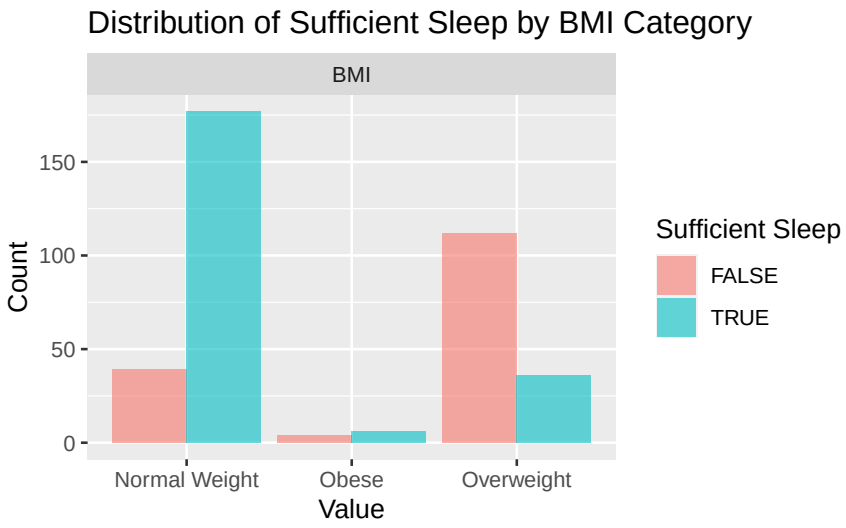
```
sleep_data %>%
  pivot_longer(cols = c(Gender), names_to = "variable", values_to = "value") %>%
  group_by(variable, value, sufficient_sleep) %>%
  summarise(count = n()) %>%
  ggplot() +
  geom_bar(
    mapping = aes(x = value, y = count, fill = sufficient_sleep),
    position = "dodge",
    alpha = 0.6,
    stat = "identity"
  ) +
```

```
  facet_wrap(~ variable, scales = "free") +
  labs(title = "Distribution of Sufficient Sleep by Gender",
       x = "Value",
       y = "Count",
       fill = "Sufficient Sleep")
```



Sleep Data BMI

```
sleep_data %>%
  pivot_longer(cols = c(BMI), names_to = "variable", values_to = "value") %>%
  mutate(value = ifelse(value == "Normal", "Normal Weight", value)) %>%
  group_by(variable, value, sufficient_sleep) %>%
  summarise(count = n()) %>%
  ggplot() +
  geom_bar(
    mapping = aes(x = value, y = count, fill = sufficient_sleep),
    position = "dodge",
    alpha = 0.6,
    stat = "identity"
  ) +
  facet_wrap(~ variable, scales = "free") +
  labs(title = "Distribution of Sufficient Sleep by BMI Category",
       x = "Value",
       y = "Count",
       fill = "Sufficient Sleep")
```



Mode

```
mode_gender <- as.character(names(which.max(table(sleep_data$Gender))))
mode_occupation <- as.character(names(which.max(table(sleep_data$Occupation))))
mode_bmi <- as.character(names(which.max(table(sleep_data$BMI))))

sleep_data <- sleep_data %>%
mutate(
  Gender = if_else(is.na(Gender), mode_gender, Gender),
  Occupation = if_else(is.na(Occupation), mode_occupation, Occupation),
  BMI = if_else(is.na(BMI), mode_bmi, BMI)
)
```

Sufficient Sleep

```
sleep_data$sufficient_sleep <- ifelse(sleep_data$Duration >= 7, "Sufficient", "Insufficient")
```

Saparate Train, Test Set

```
set.seed(123)
train_indices <- createDataPartition(sleep_data$sufficient_sleep, p = 0.7, list = FALSE)
trainingSet <- sleep_data[train_indices, ]
testSet <- sleep_data[-train_indices, ]
```

```r
trainingSet$sufficient_sleep <- as.factor(trainingSet$sufficient_sleep)
testSet$sufficient_sleep <- as.factor(testSet$sufficient_sleep)

training_Outcomes <- trainingSet$sufficient_sleep
test_Outcomes <- testSet$sufficient_sleep
```

Train

```r
model <- glm(sufficient_sleep ~ Age + Gender + Occupation + Physical + DSteps + BMI, data = tra
```

Predict

```r
predictions <- predict(model, newdata = testSet, type = "response")
```

Test

```r
threshold <- 0.5
predicted_classes <- as.factor(ifelse(predictions >= threshold, "Sufficient", "Insufficient"))
actual_classes <- test_Outcomes
accuracy <- sum(predicted_classes == actual_classes) / length(actual_classes)
print(paste("Accuracy:", accuracy))
```

```
## [1] "Accuracy: 0.981981981981982"
```

```r
model_1_preds <- testSet %>%
  add_predictions(model, type = "response") %>%
  mutate(
    outcome = as.factor(if_else(condition = pred > threshold,
                      "Sufficient", "Insufficient"))
  )
```

#Hypothesis

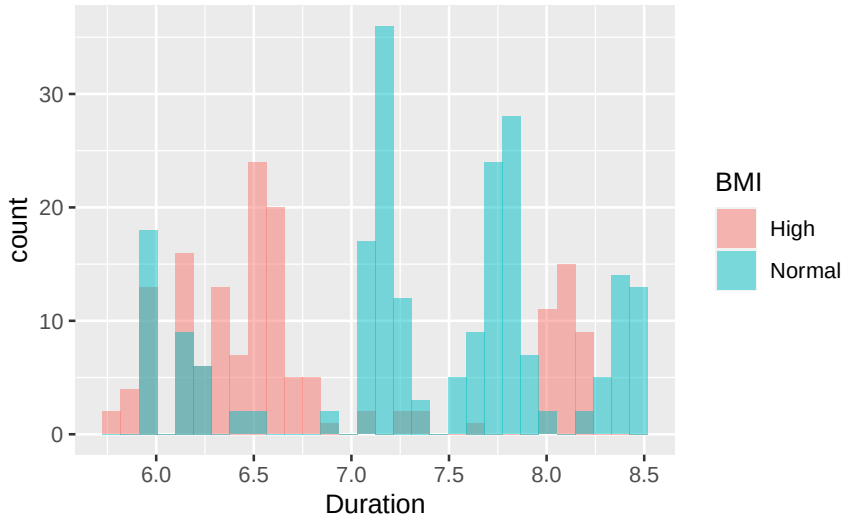
```r
Sleep_health_and_lifestyle_dataset_renamed <- Sleep_health_and_lifestyle_dataset %>%
  rename( Duration = 'Sleep Duration',
          Stress = 'Stress Level',
          Physical = 'Physical Activity Level' ,
          Quality = 'Quality of Sleep' ,
```

```
          BMI= 'BMI Category' ,
          BPressure = 'Blood Pressure' ,
          HRate = 'Heart Rate' ,
          DSteps = 'Daily Steps' ,
          Disorder = 'Sleep Disorder' )
```

```
Sleep_health_and_lifestyle_dataset_renamed$BMI[Sleep_health_and_lifestyle_dataset_renamed$BMI =
Sleep_health_and_lifestyle_dataset_renamed$BMI[Sleep_health_and_lifestyle_dataset_renamed$BMI =
Sleep_health_and_lifestyle_dataset_renamed$BMI[Sleep_health_and_lifestyle_dataset_renamed$BMI =
```

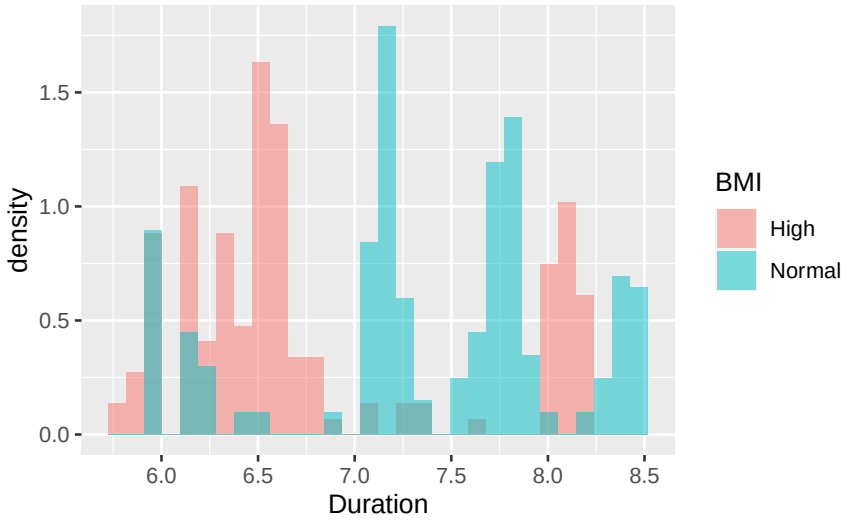
```
Sleep_health_and_lifestyle_dataset_renamed %>%
  filter(BMI == "Normal" | BMI == "High") %>%
  ggplot() +
  geom_histogram(
    mapping = aes(x = Duration, fill = BMI),
    position = "identity",
alpha = 0.5
  )
```

```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```

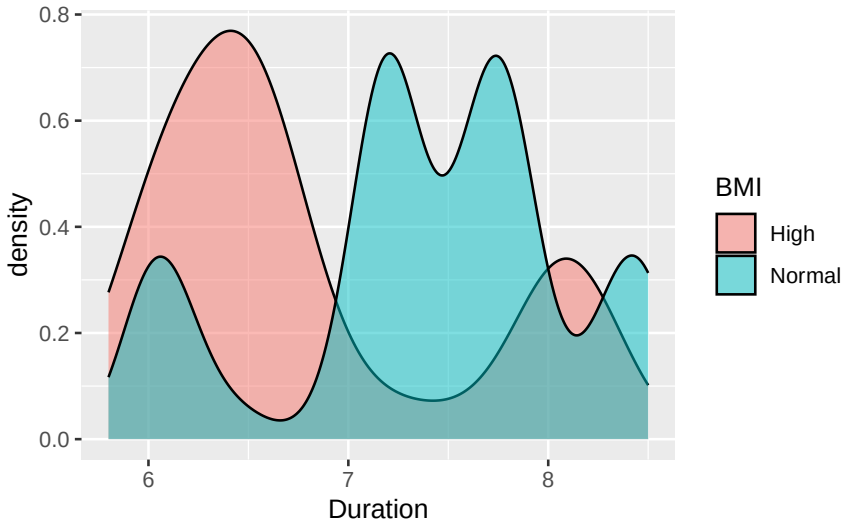



```
Sleep_health_and_lifestyle_dataset_renamed %>%
  filter(BMI == "Normal" | BMI == "High") %>%
  ggplot() +
  geom_histogram(
    mapping = aes(x = Duration, y= ..density.., fill = BMI),
    position = "identity",
alpha = 0.5
  )
```

```
## Warning: The dot-dot notation (‘..density..‘) was deprecated in ggplot2 3.4.0.
## i Please use ‘after_stat(density)‘ instead.
## This warning is displayed once every 8 hours.
## Call ‘lifecycle::last_lifecycle_warnings()‘ to see where this warning was
## generated.

## ‘stat_bin()‘ using ‘bins = 30‘. Pick better value with ‘binwidth‘.
```




```
Sleep_health_and_lifestyle_dataset_renamed %>%
  filter(BMI == "Normal" | BMI == "High") %>%
  ggplot() +
  geom_density(
    mapping = aes(x = Duration, fill = BMI),
    position = "identity",
alpha = 0.5
  )
```



```
Sleep_health_and_lifestyle_dataset_renamed %>%
summarize(
mean = mean(Duration),
median = median(Duration),
standard_deviation = sd(Duration),
minimum = min(Duration),
maximum = max(Duration)
)
```

mean	median	standard_deviation	minimum	maximum
7.132086	7.2	0.7956567	5.8	8.5

```
Model <- lm(Duration ~ BMI, data = Sleep_health_and_lifestyle_dataset_renamed)
Simulation_results <-
  Sleep_health_and_lifestyle_dataset_renamed %>%
  specify(Duration ~ BMI) %>%
  hypothesize(null = "independence") %>%
  generate(reps = 1000, type = "permute") %>%
  calculate(stat = "diff in means", order = c("Normal", "High"))
```

```
Shl_obs_stat <-
  Sleep_health_and_lifestyle_dataset_renamed %>%
  specify(formula = Duration ~ BMI) %>%
  calculate(stat = "diff in means", order = c("Normal","High"))
```

```
Shl_null <- Sleep_health_and_lifestyle_dataset_renamed %>%
  specify(Duration ~ BMI) %>%
  hypothesize(null = "independence") %>%
  generate(reps = 1000, type = "permute")
```

```
Shl_null %>%
  get_p_value(obs_stat = Shl_obs_stat, direction = "right")
```

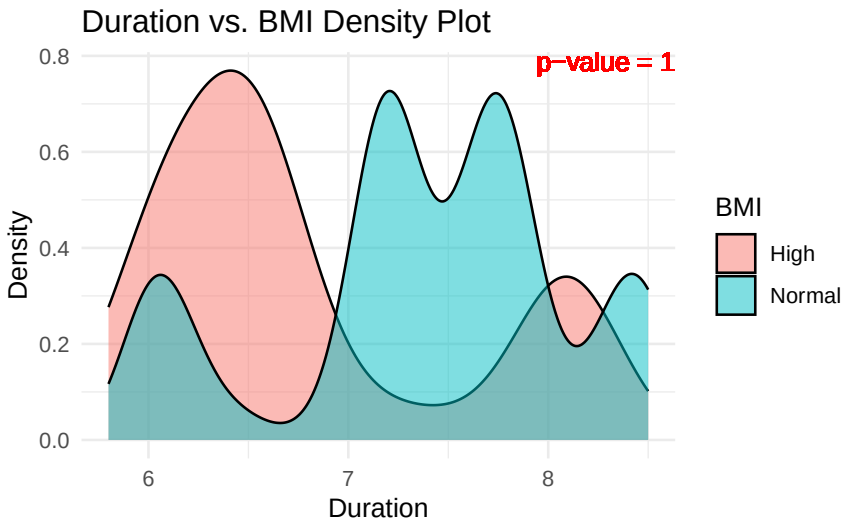
p_value
1

```
Shl_null %>% get_p_value(obs_stat = Shl_obs_stat, direction = "right")
```

p_value
1

```r
p_value <- Shl_null %>% get_p_value(obs_stat = Shl_obs_stat, direction = "right")
```

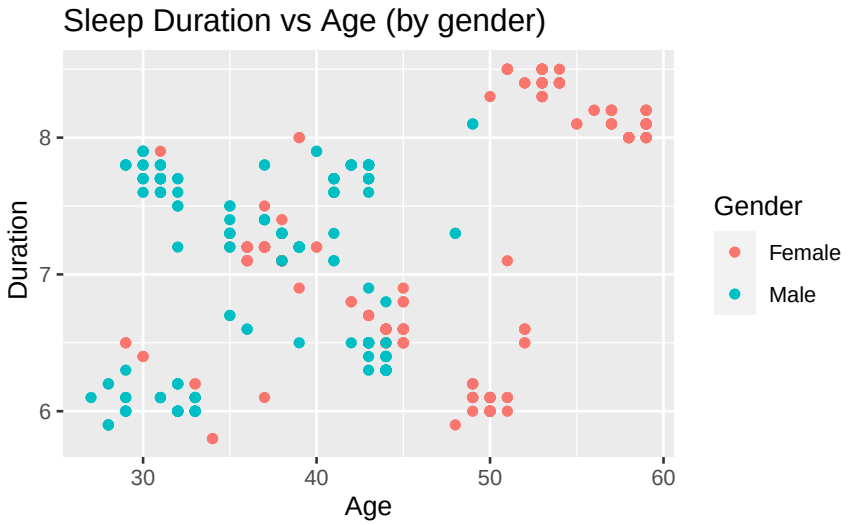
```r
Sleep_health_and_lifestyle_dataset_renamed %>%
  filter(BMI == "Normal" | BMI == "High") %>%
  ggplot() +
  geom_density(mapping = aes(x = Duration, fill = BMI), position = "identity", alpha = 0.5) +
  geom_text(aes(x = Inf, y = Inf, label = paste("p-value =", round(p_value, 3))),
            hjust = 1, vjust = 1, color = "red") +
  labs(title = "Duration vs. BMI Density Plot",
       x = "Duration",
       y = "Density") +
  theme_minimal()
```



Part 2 Data Visualization

```r
Sleep_health_and_lifestyle_dataset_renamed <- Sleep_health_and_lifestyle_dataset%>%
  rename( ID = "Person ID",
          Duration = 'Sleep Duration',
          Stress = 'Stress Level',
          Physical = 'Physical Activity Level' ,
          Quality = 'Quality of Sleep' ,
          BMI= 'BMI Category' ,
          BPressure = 'Blood Pressure' ,
          HRate = 'Heart Rate' ,
          DSteps = 'Daily Steps' ,
          Disorder = 'Sleep Disorder' )
```

```
Sleep_health_and_lifestyle_dataset_renamed %>%
  ggplot()+
  geom_point( mapping = aes( x = Age , y = Duration, color = Gender))+
  labs(
   title = "Sleep Duration vs Age (by gender)",
   x= "Age", y = " Duration")
```

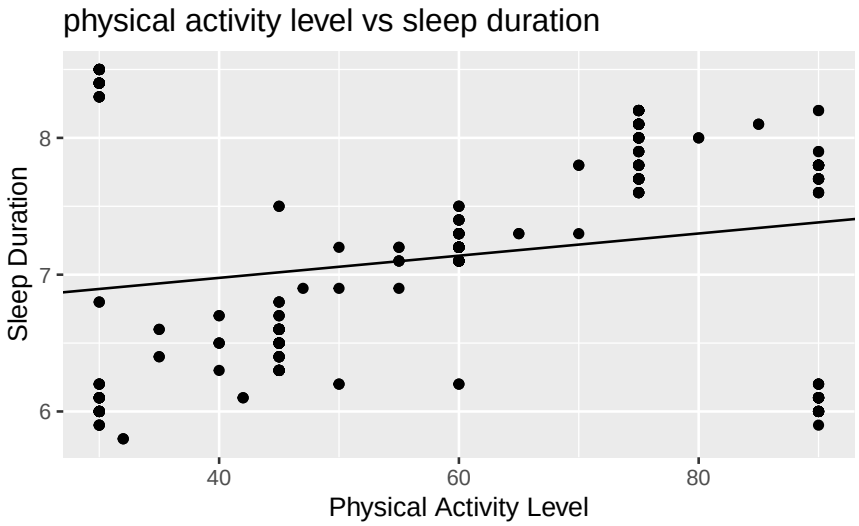



```
model_2 <- lm(Duration ~ Physical,Sleep_health_and_lifestyle_dataset_renamed)
```

```
model_2$coefficients
```

```
## (Intercept)    Physical
## 6.652127945 0.008111349
```

```
Sleep_health_and_lifestyle_dataset_renamed %>%
  ggplot() +
  geom_point(mapping = aes(x = Physical, y = Duration), bin = 10) +
  geom_abline(slope = model_2$coefficients[2],
              intercept = model_2$coefficients[1])+
   labs(x = "Physical Activity Level", y = "Sleep Duration",
                   title = "physical activity level vs sleep duration" )
```



Data Wrangling

```
Sleep_health_and_lifestyle_dataset_renamed$BMI[Sleep_health_and_lifestyle_dataset_renamed$BMI =
Sleep_health_and_lifestyle_dataset_renamed$BMI[Sleep_health_and_lifestyle_dataset_renamed$BMI =
Sleep_health_and_lifestyle_dataset_renamed$BMI[Sleep_health_and_lifestyle_dataset_renamed$BMI =
```

```
head(Sleep_health_and_lifestyle_dataset_renamed) %>%
  select(ID, HRate, Duration, Gender, Age, Occupation, Physical, BMI, Quality) %>%
  arrange(Duration)
```

ID	HRate	Duration	Gender	Age	Occupation	Physical	BMI	Quality
4	85	5.9	Male	28	Sales Representative	30	Fat	4
5	85	5.9	Male	28	Sales Representative	30	Fat	4
6	85	5.9	Male	28	Software Engineer	30	Fat	4
1	77	6.1	Male	27	Software Engineer	42	Fat	6
2	75	6.2	Male	28	Doctor	60	Normal	6
3	75	6.2	Male	28	Doctor	60	Normal	6

```
tail(Sleep_health_and_lifestyle_dataset_renamed) %>%
 select(ID, HRate, Duration, Gender, Age, Occupation, Physical, BMI, Quality) %>%
  arrange(Duration) %>%
  filter(Gender == 'Female')
```

ID	HRate	Duration	Gender	Age	Occupation	Physical	BMI	Quality
371	68	8.0	Female	59	Nurse	75	Fat	9
369	68	8.1	Female	59	Nurse	75	Fat	9

ID	HRate	Duration	Gender	Age	Occupation	Physical	BMI	Quality
370	68	8.1	Female	59	Nurse	75	Fat	9
372	68	8.1	Female	59	Nurse	75	Fat	9
373	68	8.1	Female	59	Nurse	75	Fat	9
374	68	8.1	Female	59	Nurse	75	Fat	9

```
head(Sleep_health_and_lifestyle_dataset_renamed) %>%
 select(ID, HRate, Duration, Gender, Age, Occupation, Physical, BMI, Quality) %>%
  arrange(Duration) %>%
  filter(Gender == 'Male')
```

ID	HRate	Duration	Gender	Age	Occupation	Physical	BMI	Quality
4	85	5.9	Male	28	Sales Representative	30	Fat	4
5	85	5.9	Male	28	Sales Representative	30	Fat	4
6	85	5.9	Male	28	Software Engineer	30	Fat	4
1	77	6.1	Male	27	Software Engineer	42	Fat	6
2	75	6.2	Male	28	Doctor	60	Normal	6
3	75	6.2	Male	28	Doctor	60	Normal	6

EDA

Explore dataset

```
head(Sleep_health_and_lifestyle_dataset)
```

Person ID	Gender	Age	Occupation	Sleep Duration	Quality of Sleep	Physical Activity Level	Stress Level	BMI Category	Blood Pressure	Heart Rate	Daily Steps	Sleep Disorder
1	Male	27	Software Engineer	6.1	6	42	6	Overweight	126/83	77	4200	None
2	Male	28	Doctor	6.2	6	60	8	Normal	125/80	75	10000	None
3	Male	28	Doctor	6.2	6	60	8	Normal	125/80	75	10000	None
4	Male	28	Sales Representative	5.9	4	30	8	Obese	140/90	85	3000	Sleep Apnea

Person ID	Gender	Age	Occupation	Sleep Duration	Quality of Sleep	Physical Activity Level	Stress Level	BMI Category	Blood Pressure	Heart Rate	Daily Steps	Sleep Disorder
5	Male	28	Sales Representative	5.9	4	30	8	Obese	140/90	85	3000	Sleep Apnea
6	Male	28	Software Engineer	5.9	4	30	8	Obese	140/90	85	3000	Insomnia

```
tail(Sleep_health_and_lifestyle_dataset)
```

Person ID	Gender	Age	Occupation	Sleep Duration	Quality of Sleep	Physical Activity Level	Stress Level	BMI Category	Blood Pressure	Heart Rate	Daily Steps	Sleep Disorder
369	Female	59	Nurse	8.1	9	75	3	Overweight	140/95	68	7000	Sleep Apnea
370	Female	59	Nurse	8.1	9	75	3	Overweight	140/95	68	7000	Sleep Apnea
371	Female	59	Nurse	8.0	9	75	3	Overweight	140/95	68	7000	Sleep Apnea
372	Female	59	Nurse	8.1	9	75	3	Overweight	140/95	68	7000	Sleep Apnea
373	Female	59	Nurse	8.1	9	75	3	Overweight	140/95	68	7000	Sleep Apnea
374	Female	59	Nurse	8.1	9	75	3	Overweight	140/95	68	7000	Sleep Apnea

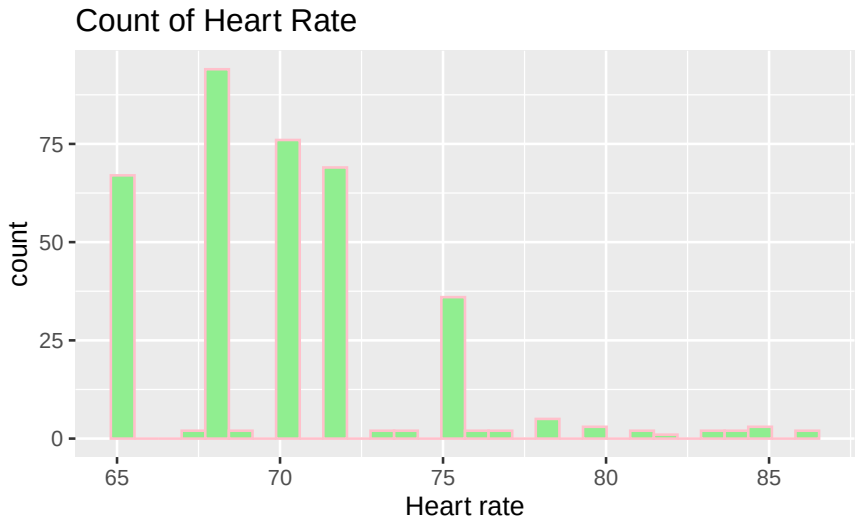
```
Sleep_health_and_lifestyle_dataset_renamed %>%
  summarize(
    standard_deviation = sd(HRate)

  )
```

standard_deviation
4.135675

Visualizing data

Histogram

```
Sleep_health_and_lifestyle_dataset_renamed %>%
  ggplot() +
    geom_histogram(mapping = aes(x = HRate), color = "pink", fill = "lightgreen") +
    labs(title = "Count of Heart Rate", x = "Heart rate")
```

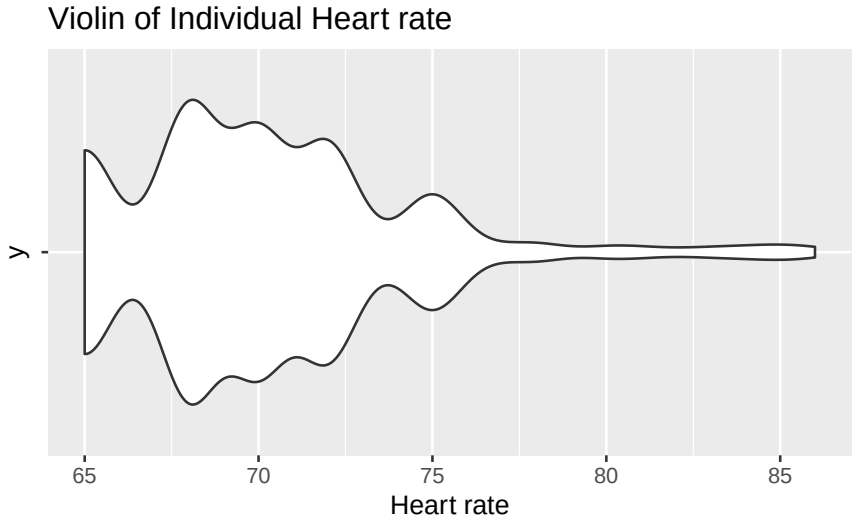


Box plot

```
Sleep_health_and_lifestyle_dataset_renamed %>%
  ggplot() +
    geom_boxplot(mapping = aes(x = HRate)) +
    labs(title = "Boxplot of Individual Heart Rate", x = "Heart rate")
```

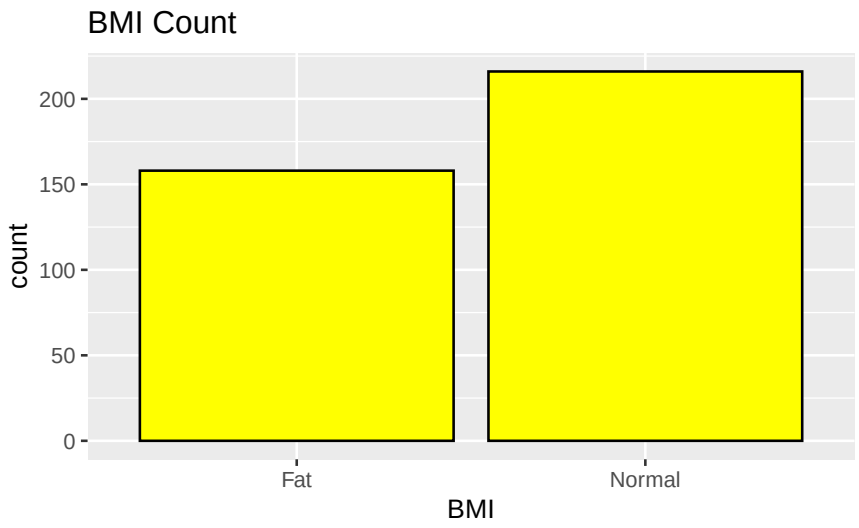
Violin plot

```
Sleep_health_and_lifestyle_dataset_renamed %>%
  ggplot() +
    geom_violin(mapping = aes(x = HRate, y ="")) +
    labs(title = "Violin of Individual Heart rate", x = "Heart rate", y = "y")
```



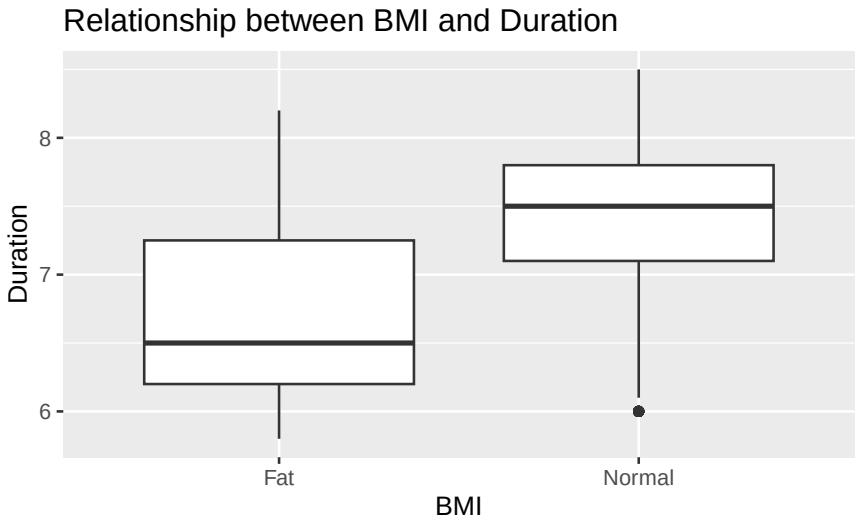
Bar Graph

```
Sleep_health_and_lifestyle_dataset_renamed %>%
  ggplot() +
    geom_bar(mapping = aes(x = BMI), color = "black", fill = "yellow") +
    labs(title = "BMI Count", x = "BMI")
```



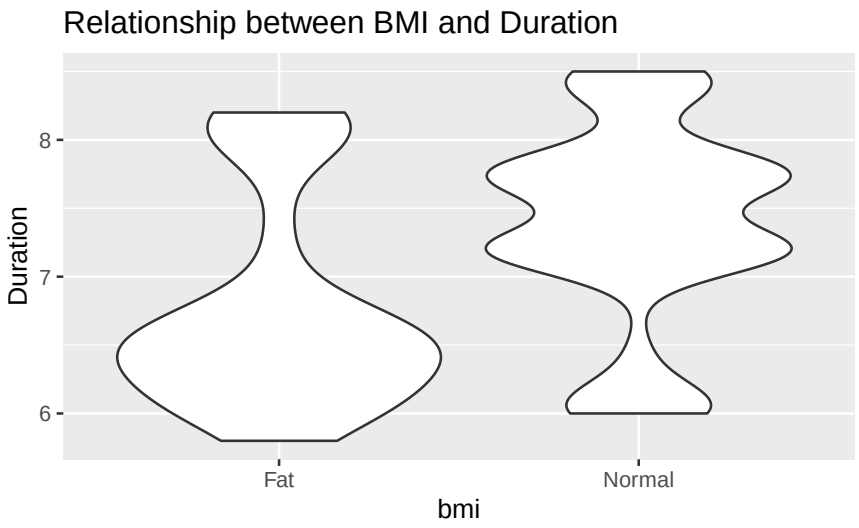
Box plot

```
Sleep_health_and_lifestyle_dataset_renamed %>%
  ggplot() +
    geom_boxplot(mapping = aes(x = BMI, y = Duration)) +
    labs(title = "Relationship between BMI and Duration", x = "BMI")
```



Violin plot

```
Sleep_health_and_lifestyle_dataset_renamed %>%
  ggplot() +
    geom_violin(mapping = aes(x = BMI, y = Duration)) +
    labs(title = "Relationship between BMI and Duration", x = "bmi", y = "Duration")
```

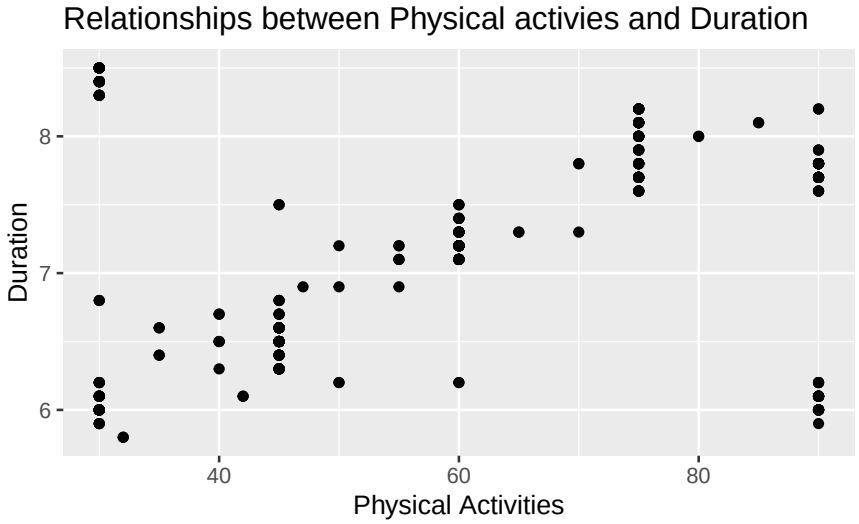


#Scatter plot_Duration and Heart Rate

```
Sleep_health_and_lifestyle_dataset_renamed    %>%
ggplot()    +
geom_point(mapping = aes(x = BMI, y = Duration))    +
labs(
title = "Scatter plot of Duration and Heart Rate",
x = "BMI",
y = "Duration"
)
```

PART 5 _ Modeling

```r
Sleep_health_and_lifestyle_dataset_renamed%>%
  ggplot()+
  geom_point( mapping = aes( x  = Physical , y = Duration)) +
  labs(title = "Relationships between Physical activies and Duration",
       x = "Physical Activities" , y = "Duration")
```

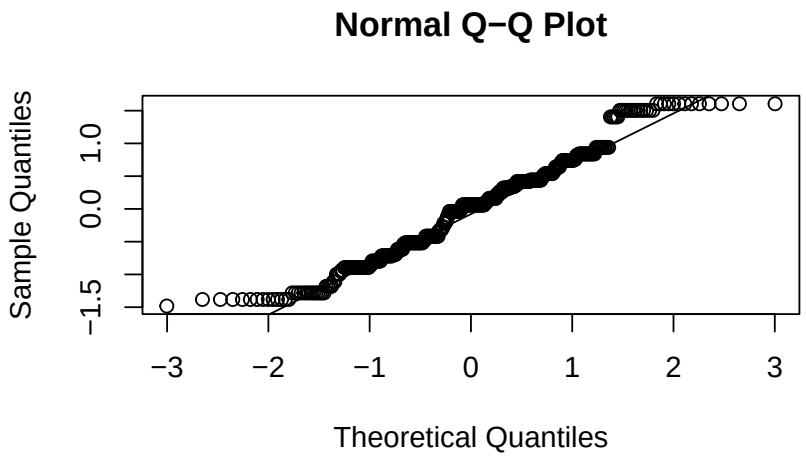



```r
data <- Sleep_health_and_lifestyle_dataset_renamed

model <- lm(Duration ~ Physical, data = Sleep_health_and_lifestyle_dataset_renamed)


summary(model)
```

```
##
## Call:
## lm(formula = Duration ~ Physical, data = Sleep_health_and_lifestyle_dataset_renamed)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -1.48215 -0.59686  0.06119  0.43952  1.60453
##
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)
## (Intercept) 6.652128   0.121379  54.805  < 2e-16 ***
## Physical    0.008111   0.001935   4.191 3.47e-05 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.7786 on 372 degrees of freedom
## Multiple R-squared:  0.0451, Adjusted R-squared:  0.04253
```

```
## F-statistic: 17.57 on 1 and 372 DF,  p-value: 3.467e-05
```

```
residuals <- residuals(model)

qqnorm(residuals)
qqline(residuals)
```




```
labs( title  = "QQplot" , x = "Theoretical" , y = "Quantaties")
```

```
## $x
## [1] "Theoretical"
##
## $y
## [1] "Quantaties"
##
## $title
## [1] "QQplot"
##
## attr(,"class")
## [1] "labels"
```

```
Renamed_other_model <- lm(Duration ~ Physical, data = Sleep_health_and_lifestyle_dataset_renam
```

```
Renamed_other_model$coefficients
```

```
## (Intercept)    Physical
## 6.652127945 0.008111349
```

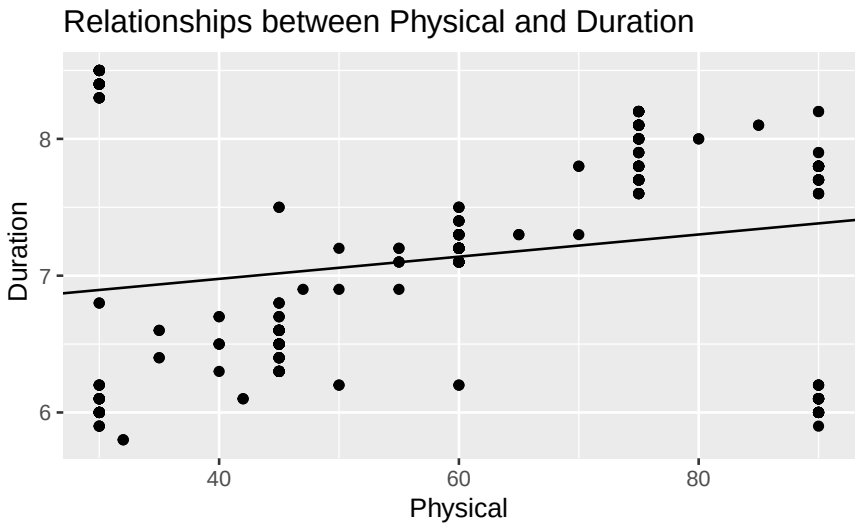
```
Renamed_other_model%>%
  tidy()
```

term	estimate	std.error	statistic	p.value
(Intercept)	6.6521279	0.1213792	54.804523	0.00e+00
Physical	0.0081113	0.0019352	4.191459	3.47e-05

```
Renamed_other_model%>%
  glance()%>%
  select(r.squared)
```

r.squared
0.0450969

```
Sleep_health_and_lifestyle_dataset_renamed%>%
  ggplot()+
  geom_point(mapping = aes( x  = Physical , y = Duration) )+
  geom_abline(slope = Renamed_other_model$coefficients[2]   ,
              intercept = Renamed_other_model$coefficients[1]  )+
  labs( title = "Relationships between Physical and Duration",
        x = " Physical ",
        y = " Duration" )
```



#Advanced Modeling

```
continuous_model <- lm(Duration ~ Gender + Age + Occupation + DSteps + BMI + Physical, data = S
coefficients <- tidy (continuous_model)
coefficients
```

term	estimate	std.error	statistic	p.value
(Intercept)	3.8296315	0.3275001	11.6935269	0.0000000
GenderMale	-0.1949526	0.1287189	-1.5145603	0.1307662
Age	0.0655149	0.0062353	10.5071674	0.0000000
OccupationDoctor	0.4245451	0.1384180	3.0671237	0.0023258
OccupationEngineer	0.3033566	0.1400149	2.1666029	0.0309244
OccupationLawyer	0.2877110	0.1525615	1.8858691	0.0601220
OccupationManager	0.1297888	0.4617227	0.2810969	0.7787985
OccupationNurse	-0.2098419	0.1178946	-1.7799108	0.0759387
OccupationSales Representative	0.4350443	0.3563943	1.2206825	0.2230096
OccupationSalesperson	0.3071433	0.1756392	1.7487178	0.0811969
OccupationScientist	0.3016381	0.2640060	1.1425423	0.2539922
OccupationSoftware Engineer	0.7486436	0.2586419	2.8945175	0.0040302
OccupationTeacher	0.3402722	0.1224097	2.7797809	0.0057268
DSteps	-0.0002563	0.0000246	-10.4369782	0.0000000
BMINormal	1.1644995	0.1087480	10.7082352	0.0000000
Physical	0.0254908	0.0020941	12.1729446	0.0000000

```
r_squared <- glance(continuous_model)$r.squared
```

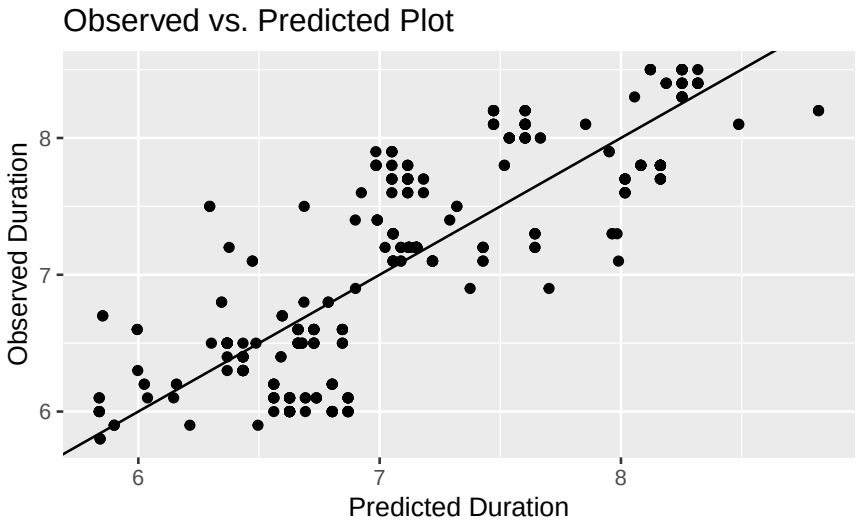
```
Sleep_health_and_lifestyle_dataset_df <- Sleep_health_and_lifestyle_dataset_renamed %>%
  add_predictions(continuous_model) %>%
  add_residuals(continuous_model)
```

#Histogram of residual in Sleep_health_and_lifestyle_dataset_df

```
Sleep_health_and_lifestyle_dataset_df %>%
  ggplot() +
  geom_histogram(mapping = aes(x = resid), color = "blue", fill = "pink", bins = 10) +
  labs(x = "residual", y ="Duration",
title = "Histogram of residual in Sleep_health_and_lifestyle_dataset_df")
```

#Observed vs. Predicted Plot

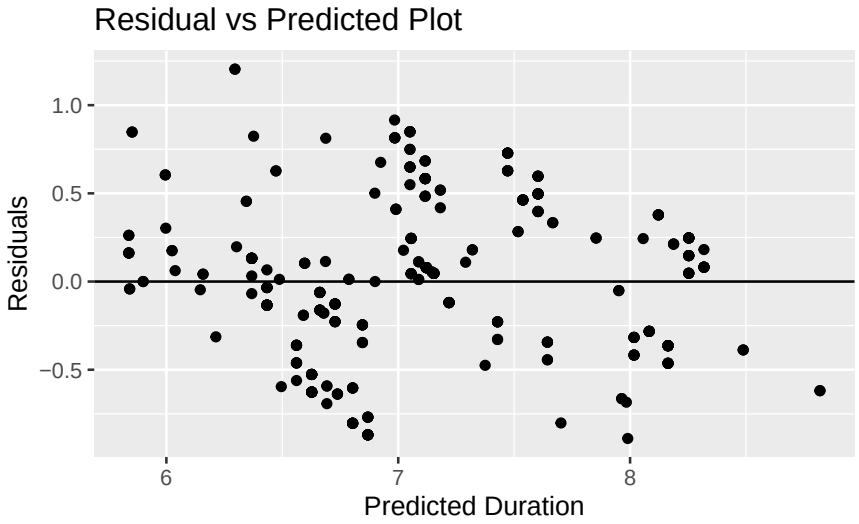
```
Sleep_health_and_lifestyle_dataset_df %>%
  ggplot() +
  geom_point(mapping = aes(x = pred, y = Duration)) +
  geom_abline(slope = 1, intercept = 0) +
  labs(title = "Observed vs. Predicted Plot", x = "Predicted Duration", y = "Observed Duration"
```



#Residual vs Predicted Plot

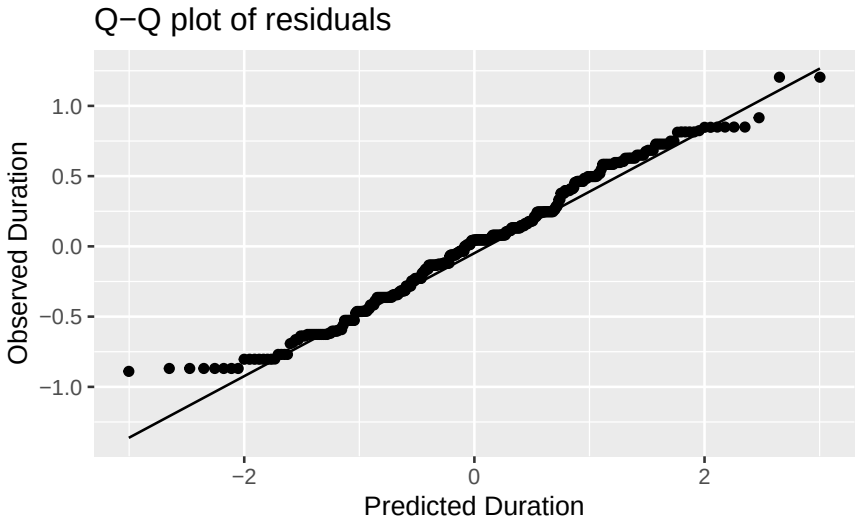
```
Sleep_health_and_lifestyle_dataset_df %>%
  ggplot() +
  geom_point(mapping = aes(x = pred, y = resid)) +
  geom_hline(yintercept = 0) +
```

```
  labs( title= "Residual vs Predicted Plot",
        x = "Predicted Duration",
        y = "Residuals")
```



#Q-Q Plot (Obeserved vs Predicted Plot)

```
Sleep_health_and_lifestyle_dataset_df %>%
  ggplot() +
  geom_qq(aes(sample = resid)) +
  geom_qq_line(aes(sample = resid))+
  labs(title = "Q-Q plot of residuals", x= "Predicted Duration", y= "Observed Duration")
```



#Box Plot

```r
Sleep_health_and_lifestyle_dataset_df %>%
  pivot_longer(
    cols = Gender:Occupation | Physical | BMI | DSteps,
    names_to = "column",
    values_to = "value",
    values_transform = list(value = 'factor')
) %>%
ggplot() +
  geom_boxplot(aes(x = reorder(value, Duration, FUN = median), y = Duration)) +
  facet_wrap(~column, scales = "free_x") +
  labs(x = "x variable", y = "Duration", title = "Box Plot of x variables") +
  theme(axis.text.x = element_text(angle = 45))
```