

# week.5..p

sahrash fatima Lab

2024-10-02

```
# Load the dataset from the specified path
dataset <- read.csv("D:/iupui/2rd sem/R- stats/parkinsons_disease_data.csv")
```

```
# View the first few rows of the dataset to verify it loaded correctly
head(dataset)
```

```
## PatientID Age Gender Ethnicity EducationLevel BMI Smoking
## 1 3058 85 0 3 1 19.61988 0
## 2 3059 75 0 0 2 16.24734 1
## 3 3060 70 1 0 0 15.36824 0
## 4 3061 52 0 0 0 15.45456 0
## 5 3062 87 0 0 1 18.61604 0
## 6 3063 68 1 2 1 39.42331 1
## AlcoholConsumption PhysicalActivity DietQuality SleepQuality
## 1 5.108241 1.3806599 3.893969 9.283194
## 2 6.027648 8.4098041 8.513428 5.602470
## 3 2.242135 0.2132746 6.498805 9.929824
## 4 5.997788 1.3750452 6.715033 4.196189
## 5 9.775243 1.1886071 4.657572 9.363925
## 6 13.596889 7.7967040 7.070239 7.737549
## FamilyHistoryParkinsons TraumaticBrainInjury Hypertension Diabetes
Depression
## 1 0 0 0 0
0
## 2 0 0 0 0
0
## 3 0 0 0 1
0
## 4 0 0 0 0
0
## 5 0 0 0 0
0
## 6 0 0 0 0
0
## Stroke SystolicBP DiastolicBP CholesterolTotal CholesterolLDL
CholesterolHDL
## 1 0 129 60 222.8423 148.12562
37.86778
## 2 0 163 76 210.5011 153.75646
77.22812
## 3 0 113 93 287.3880 118.70260
85.58830
## 4 0 146 78 280.3395 136.29919
```

```

51.86963
## 5      0      115      94      284.0142      108.44945
25.06942
## 6      0      151      90      290.1331      91.75022
54.48892
## CholesterolTriglycerides      UPDRS      MoCA FunctionalAssessment
Tremor
## 1      337.3071      6.458713      29.181289      1.572427
1
## 2      264.6355      37.306703      12.332639      4.787551
0
## 3      395.6626      67.838170      29.927783      2.130686
1
## 4      362.1897      52.964696      21.304268      3.391288
1
## 5      149.9566      21.804880      8.336364      3.200969
0
## 6      253.7973      101.912536      27.370580      6.824779
0
## Rigidity Bradykinesia PosturalInstability SpeechProblems SleepDisorders
## 1      0      0      0      0      0
## 2      1      0      1      0      1
## 3      0      0      0      1      0
## 4      1      1      0      0      0
## 5      0      0      1      0      1
## 6      0      0      0      0      0
## Constipation Diagnosis DoctorInCharge
## 1      0      0      DrXXXConfid
## 2      0      1      DrXXXConfid
## 3      1      1      DrXXXConfid
## 4      1      1      DrXXXConfid
## 5      0      0      DrXXXConfid
## 6      0      0      DrXXXConfid

```

*# Check the structure of the dataset to ensure variables are correctly loaded*  
**str(dataset)**

```

## 'data.frame':    2105 obs. of  35 variables:
## $ PatientID      : int  3058 3059 3060 3061 3062 3063 3064 3065
3066 3067 ...
## $ Age            : int  85 75 70 52 87 68 78 70 80 71 ...
## $ Gender         : int  0 0 1 0 0 1 1 1 0 0 ...
## $ Ethnicity      : int  3 0 0 0 0 2 0 0 2 3 ...
## $ EducationLevel : int  1 2 0 0 1 1 0 0 1 2 ...
## $ BMI            : num  19.6 16.2 15.4 15.5 18.6 ...
## $ Smoking        : int  0 1 0 0 0 1 1 1 1 1 ...
## $ AlcoholConsumption : num  5.11 6.03 2.24 6 9.78 ...
## $ PhysicalActivity : num  1.381 8.41 0.213 1.375 1.189 ...
## $ DietQuality     : num  3.89 8.51 6.5 6.72 4.66 ...
## $ SleepQuality    : num  9.28 5.6 9.93 4.2 9.36 ...

```

```
## $ FamilyHistoryParkinsons : int 0 0 0 0 0 0 0 0 0 0 ...
## $ TraumaticBrainInjury    : int 0 0 0 0 0 0 0 0 0 1 ...
## $ Hypertension            : int 0 0 0 0 0 0 1 0 0 0 ...
## $ Diabetes                : int 0 0 1 0 0 0 0 1 1 0 ...
## $ Depression              : int 0 0 0 0 0 0 0 0 0 1 ...
## $ Stroke                  : int 0 0 0 0 0 0 0 0 0 0 ...
## $ SystolicBP              : int 129 163 113 146 115 151 122 129 133 169
...
## $ DiastolicBP             : int 60 76 93 78 94 90 60 99 113 105 ...
## $ CholesterolTotal        : num 223 211 287 280 284 ...
## $ CholesterolLDL          : num 148 154 119 136 108 ...
## $ CholesterolHDL          : num 37.9 77.2 85.6 51.9 25.1 ...
## $ CholesterolTriglycerides: num 337 265 396 362 150 ...
## $ UPDRS                   : num 6.46 37.31 67.84 52.96 21.8 ...
## $ MoCA                    : num 29.18 12.33 29.93 21.3 8.34 ...
## $ FunctionalAssessment    : num 1.57 4.79 2.13 3.39 3.2 ...
## $ Tremor                  : int 1 0 1 1 0 0 1 1 0 0 ...
## $ Rigidity                 : int 0 1 0 1 0 0 0 0 0 0 ...
## $ Bradykinesia            : int 0 0 0 1 0 0 0 0 0 0 ...
## $ PosturalInstability     : int 0 1 0 0 1 0 0 1 0 0 ...
## $ SpeechProblems          : int 0 0 1 0 0 0 1 0 0 0 ...
## $ SleepDisorders          : int 0 1 0 0 1 0 0 0 0 1 ...
## $ Constipation            : int 0 0 1 1 0 0 0 1 0 0 ...
## $ Diagnosis               : int 0 1 1 1 0 0 0 1 1 0 ...
## $ DoctorInCharge          : chr "DrXXXConfid" "DrXXXConfid"
"DrXXXConfid" "DrXXXConfid" ...
```

*# Install and load required packages*

```
if (!require(ggplot2)) install.packages("ggplot2", dependencies=TRUE)
```

```
## Loading required package: ggplot2
```

```
if (!require(psych)) install.packages("psych", dependencies=TRUE)
```

```
## Loading required package: psych
```

```
##
```

```
## Attaching package: 'psych'
```

```
## The following objects are masked from 'package:ggplot2':
```

```
##
```

```
##      %+%, alpha
```

```
if (!require(corrplot)) install.packages("corrplot", dependencies=TRUE)
```

```
## Loading required package: corrplot
```

```
## corrplot 0.94 loaded
```

```
library(ggplot2)
```

```
library(psych)
```

```
library(corrplot)
```

```

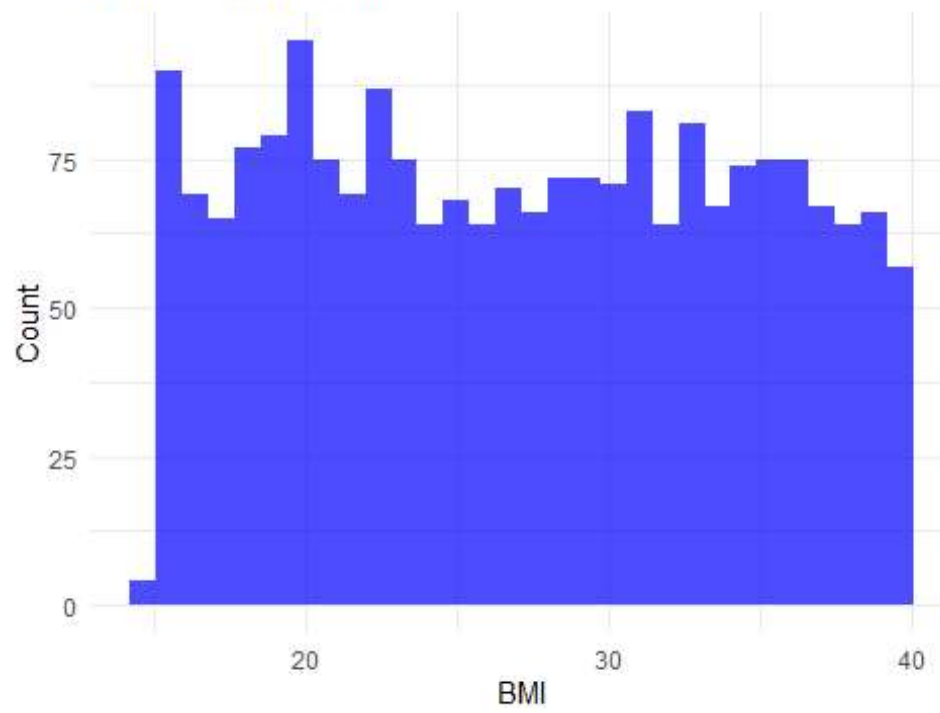
# Plot distribution of variables and check for skewness
plot_distribution <- function(variable_name, dataset) {
  tryCatch({
    ggplot(dataset, aes(x = !!sym(variable_name))) +
      geom_histogram(bins = 30, fill = "blue", alpha = 0.7) +
      labs(title = paste("Distribution of", variable_name), x =
variable_name, y = "Count") +
      theme_minimal()
  }, error = function(e) {
    return(paste("Error in plotting", variable_name, ":", e$message))
  })
}

# Variables to plot (based on dataset)
variables <- c("BMI", "UPDRS", "MoCA", "PhysicalActivity", "SystolicBP")

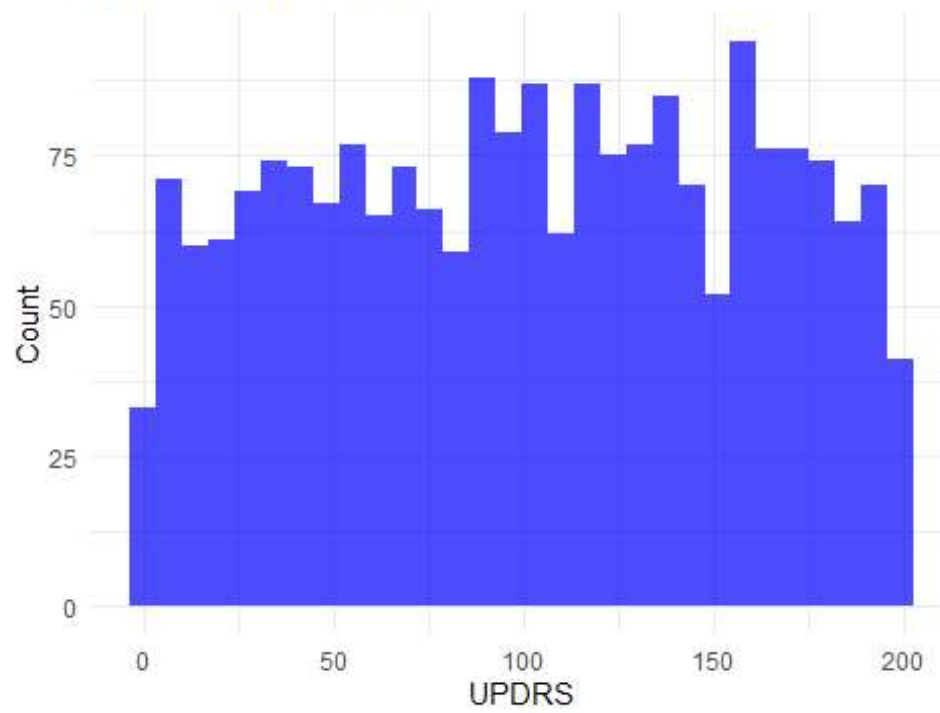
# Loop through and plot distributions
for (var in variables) {
  print(plot_distribution(var, dataset))
}

```

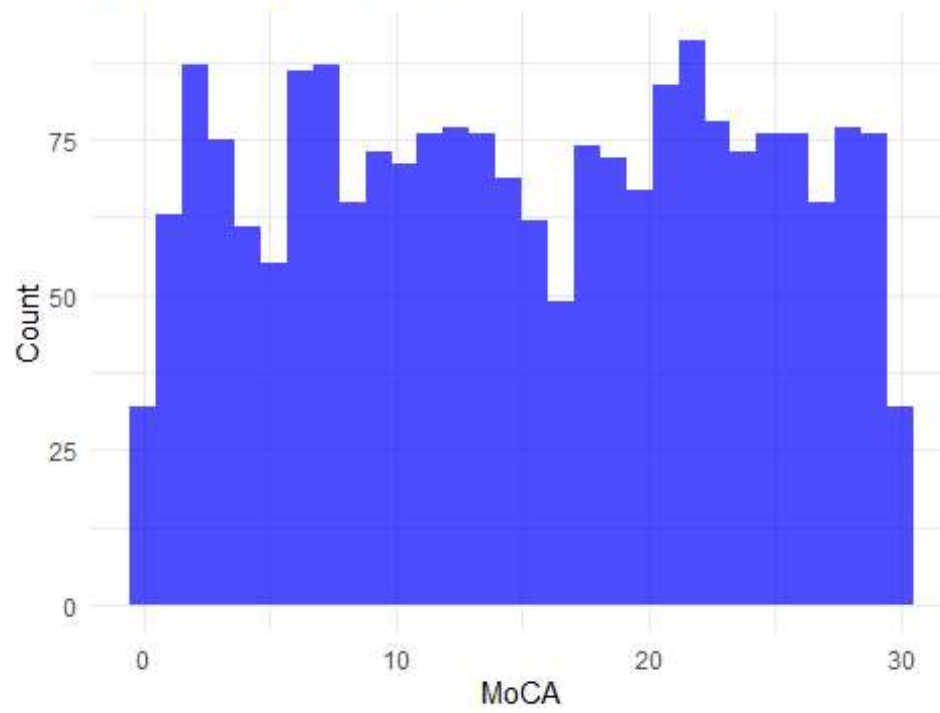
Distribution of BMI



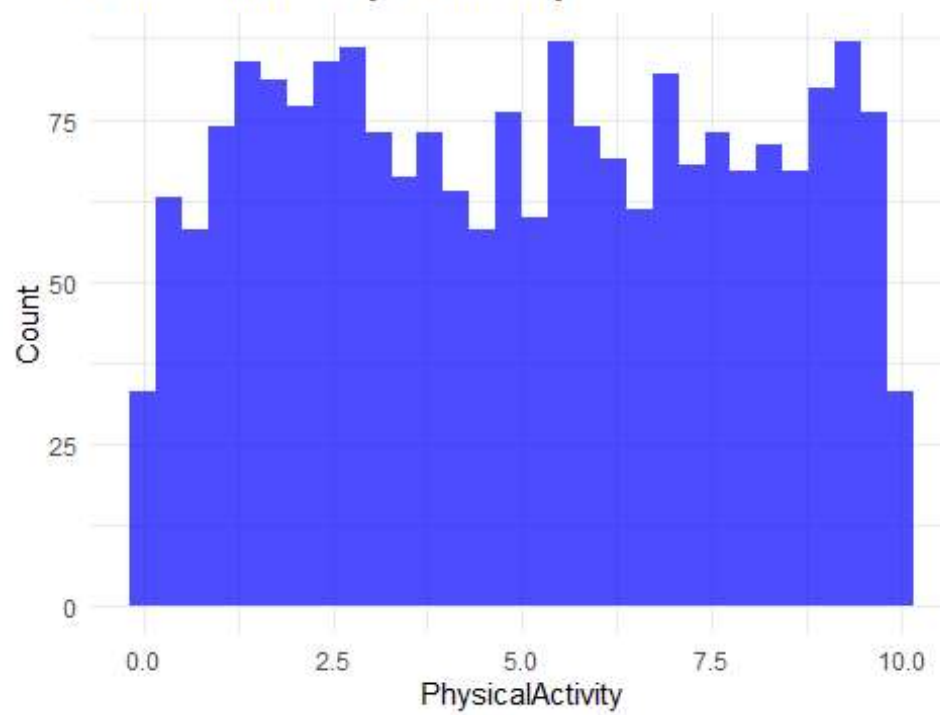
Distribution of UPDRS

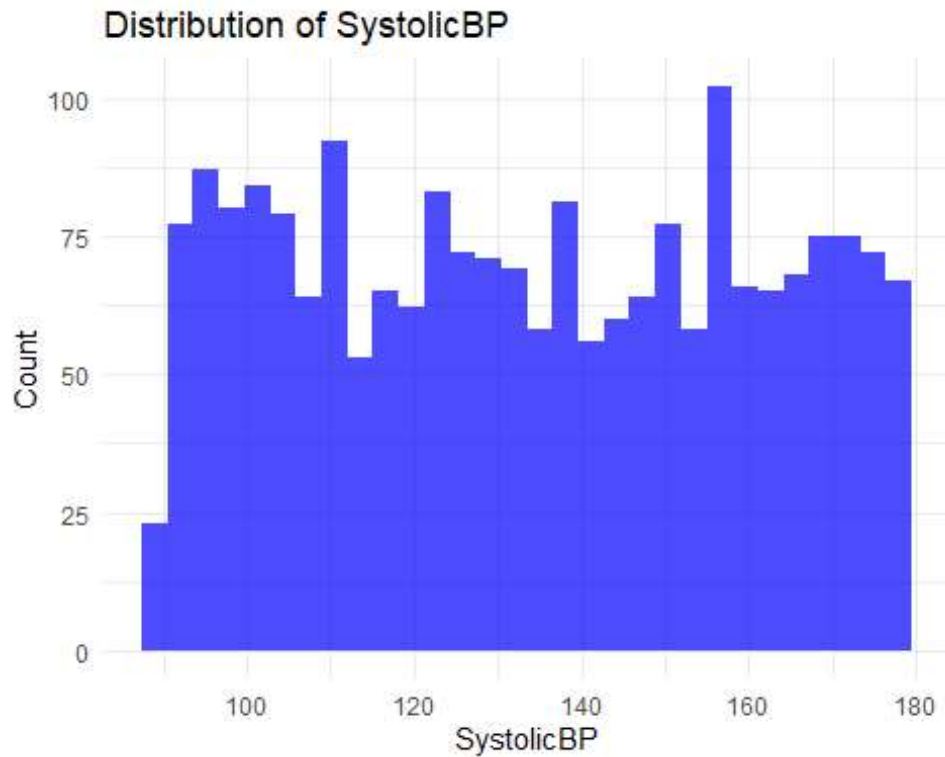


Distribution of MoCA



Distribution of PhysicalActivity





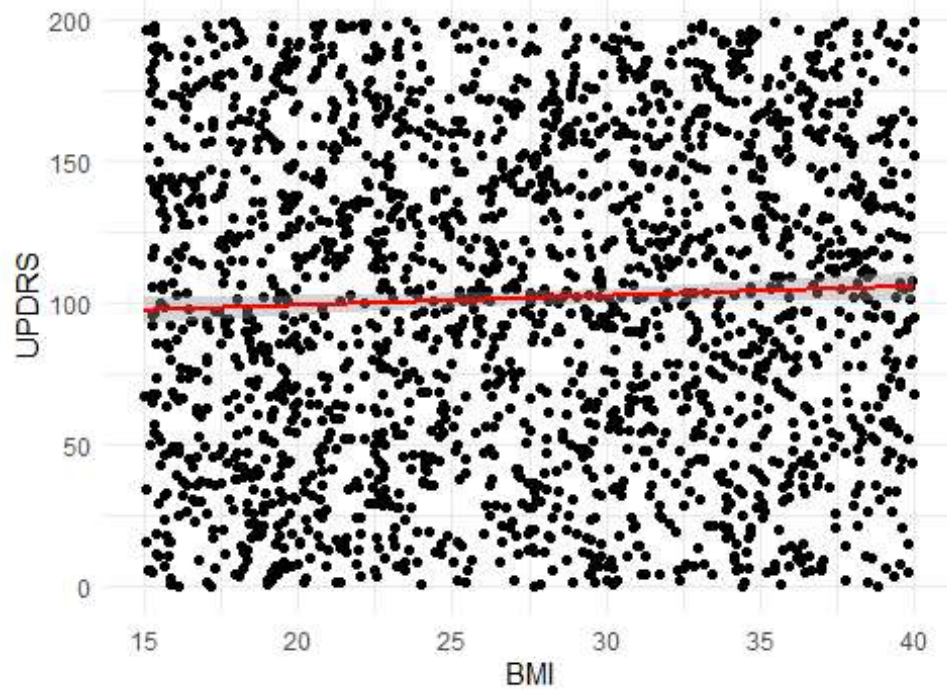
```
# Scatter plot function with regression line
scatter_plot_with_regression <- function(predictor, outcome, dataset) {
  tryCatch({
    ggplot(dataset, aes(x = !!sym(predictor), y = !!sym(outcome))) +
      geom_point() +
      geom_smooth(method = "lm", color = "red") +
      labs(title = paste("Scatter Plot of", predictor, "vs", outcome),
           x = predictor, y = outcome) +
      theme_minimal()
  }, error = function(e) {
    return(paste("Error in plotting", predictor, "vs", outcome, ":",
e$message))
  })
}

# Variables to plot against UPDRS (adjust if needed)
predictors <- c("BMI", "MoCA", "PhysicalActivity", "SystolicBP")
outcome <- "UPDRS"

# Loop through and plot
for (var in predictors) {
  print(scatter_plot_with_regression(var, outcome, dataset))
}

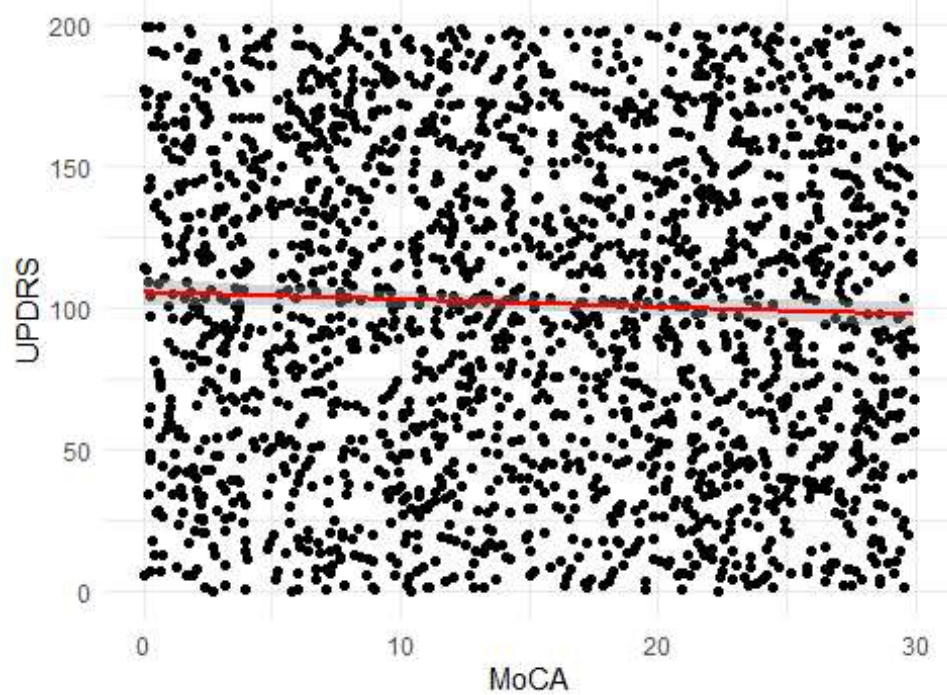
## `geom_smooth()` using formula = 'y ~ x'
```

Scatter Plot of BMI vs UPDRS



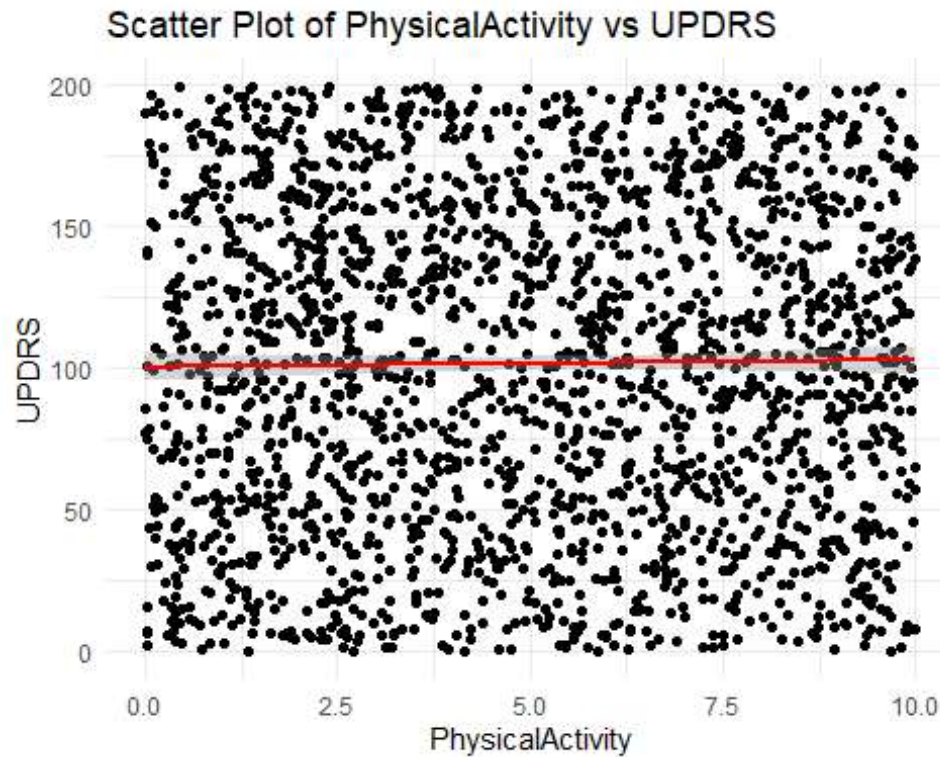
```
## `geom_smooth()` using formula = 'y ~ x'
```

Scatter Plot of MoCA vs UPDRS

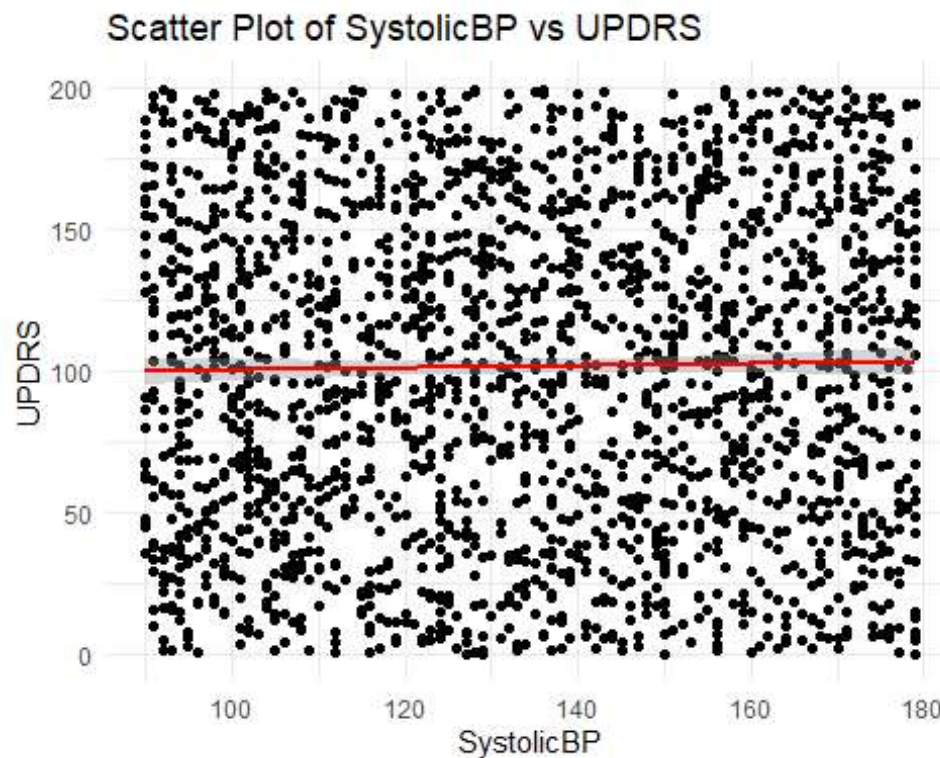


```
## `geom_smooth()` using formula = 'y ~ x'
```





```
## `geom_smooth()` using formula = 'y ~ x'
```



''' Correlations:

Because the regression lines in all of the scatter plots are almost flat, it appears that none of the variables (BMI, SystolicBP, PhysicalActivity, MoCA, and Physical Activity)

have a significant linear association with UPDRS. Distributions: Each variable's data is approximately uniform, with no significant skewness or strong trends in any direction, according to the histograms for the distributions.