

# Untitled

2024-05-05

```
library(readxl)
riseholme_data_24_1_ <- read_excel("C:/Users/Lenovo/Desktop/R/riseholme_data_24(1).xlsx")
View(riseholme_data_24_1_)
```

```
library(readxl)
library(dplyr)
```

```
##
## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':
##
##   filter, lag

## The following objects are masked from 'package:base':
##
##   intersect, setdiff, setequal, union
```

```
library(broom)
```

```
str(riseholme_data_24_1_)
```

```
## tibble [54 x 12] (S3: tbl_df/tbl/data.frame)
##  $ Group_number   : chr [1:54] "G1" "G1" "G1" "G1" ...
##  $ site           : chr [1:54] "old_wood" "old_wood" "old_wood" "old_wood" ...
##  $ X_location      : num [1:54] 2 9 15 12 4 4 21 30 18 9 ...
##  $ Y_location      : num [1:54] 6 4 5 20 18 4 21 30 0 4 ...
##  $ tree_DBH        : num [1:54] 3 55 33 20 36 12 20 19 3 12 ...
##  $ neigh_5m        : num [1:54] 17 11 11 100 34 3 6 7 7 2 ...
##  $ max_neigh_dbh    : num [1:54] 55 19 21 12 4 19 58 24 16 13.5 ...
##  $ ground_cover     : num [1:54] 30 35 1 25 15 35 60 5 80 70 ...
##  $ number_flowers   : num [1:54] 0 2 0 4 0 0 0 0 0 0 ...
##  $ num_diff_colour  : num [1:54] 0 2 0 2 0 0 0 0 0 0 ...
##  $ radius           : num [1:54] 5 5 5 5 5 5 5 5 5 5 ...
##  $ n_neigh_m2       : num [1:54] 0.216 0.14 0.14 1.273 0.433 ...
```

```
summary(riseholme_data_24_1_)
```

```
##  Group_number      site      X_location      Y_location
## Length:54      Length:54      Min.   : 0.00      Min.   : 0.00
## Class :character Class :character 1st Qu.: 7.00      1st Qu.: 9.00
```

```
## Mode :character Mode :character Median :11.50 Median :18.00
## Mean :13.76 Mean :15.85
## 3rd Qu.:20.50 3rd Qu.:23.00
## Max. :30.00 Max. :30.00
## tree_DBH neigh_5m max_neigh_dbh ground_cover
## Min. : 1.00 Min. : 2.00 Min. : 3.00 Min. : 1.0
## 1st Qu.: 11.25 1st Qu.: 6.00 1st Qu.: 12.25 1st Qu.: 7.0
## Median : 16.00 Median : 9.00 Median : 17.50 Median :17.0
## Mean : 25.80 Mean : 20.37 Mean : 22.90 Mean :25.5
## 3rd Qu.: 37.50 3rd Qu.: 28.25 3rd Qu.: 28.50 3rd Qu.:35.0
## Max. :167.00 Max. :113.00 Max. :128.00 Max. :97.0
## number_flowers num_diff_colour radius n_neigh_m2
## Min. : 0.000 Min. :0.0 Min. :3.000 Min. :0.02546
## 1st Qu.: 0.000 1st Qu.:0.0 1st Qu.:5.000 1st Qu.:0.07639
## Median : 0.000 Median :0.0 Median :5.000 Median :0.13369
## Mean : 8.593 Mean :0.5 Mean :4.815 Mean :0.27445
## 3rd Qu.: 3.500 3rd Qu.:1.0 3rd Qu.:5.000 3rd Qu.:0.36535
## Max. :117.000 Max. :2.0 Max. :5.000 Max. :1.43876
```

```
library(tibble)
library(ggplot2)
```

```
model <- lm(tree_DBH ~ neigh_5m + max_neigh_dbh + ground_cover, data = riseholme_data_24_1_)
```

```
summary_model <- summary(model)
```

```
coefficients <- summary_model$coefficients
coefficients
```

```
## Estimate Std. Error t value Pr(>|t|)
## (Intercept) 12.3202667 6.0169065 2.0476082 0.045869747
## neigh_5m 0.3823110 0.1285439 2.9741663 0.004513278
## max_neigh_dbh -0.1090730 0.1650639 -0.6607929 0.511778376
## ground_cover 0.3210111 0.1396511 2.2986645 0.025740839
```

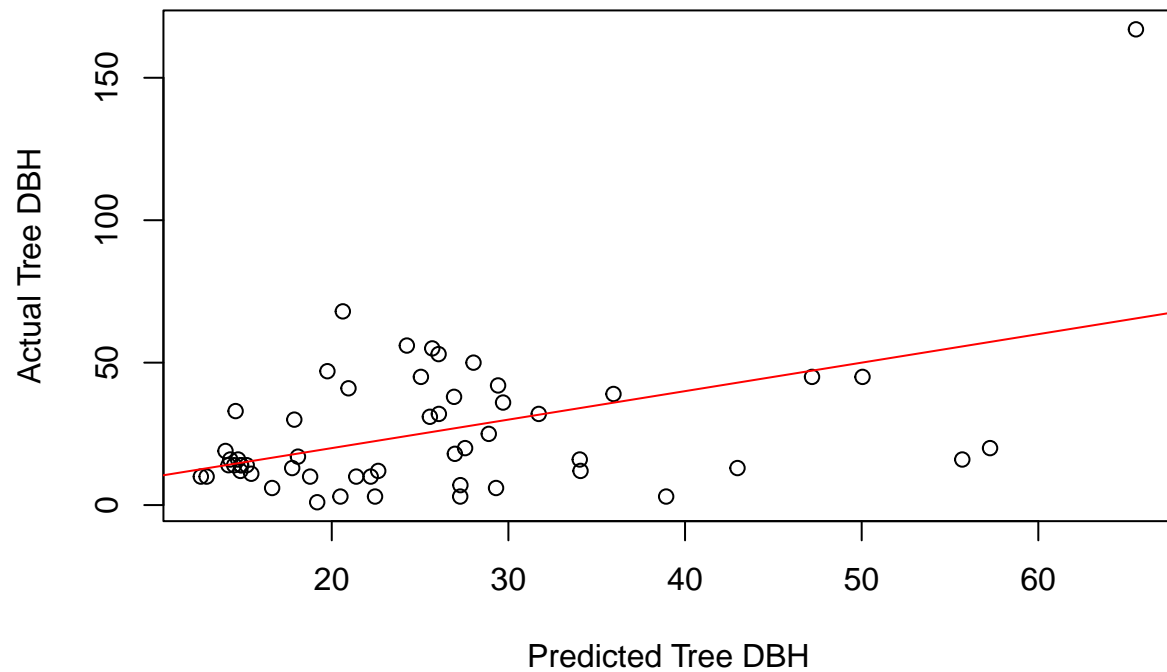
```
# Get predicted values from the model
```

```
riseholme_data_24_1_$predicted_tree_DBH <- predict(model)
```

```
# Scatter plot of actual vs predicted tree_DBH
```

```
plot(riseholme_data_24_1_$predicted_tree_DBH, riseholme_data_24_1_$tree_DBH,
     xlab = "Predicted Tree DBH", ylab = "Actual Tree DBH",
     main = "Actual vs Predicted Tree DBH")
abline(0, 1, col = "red")
```

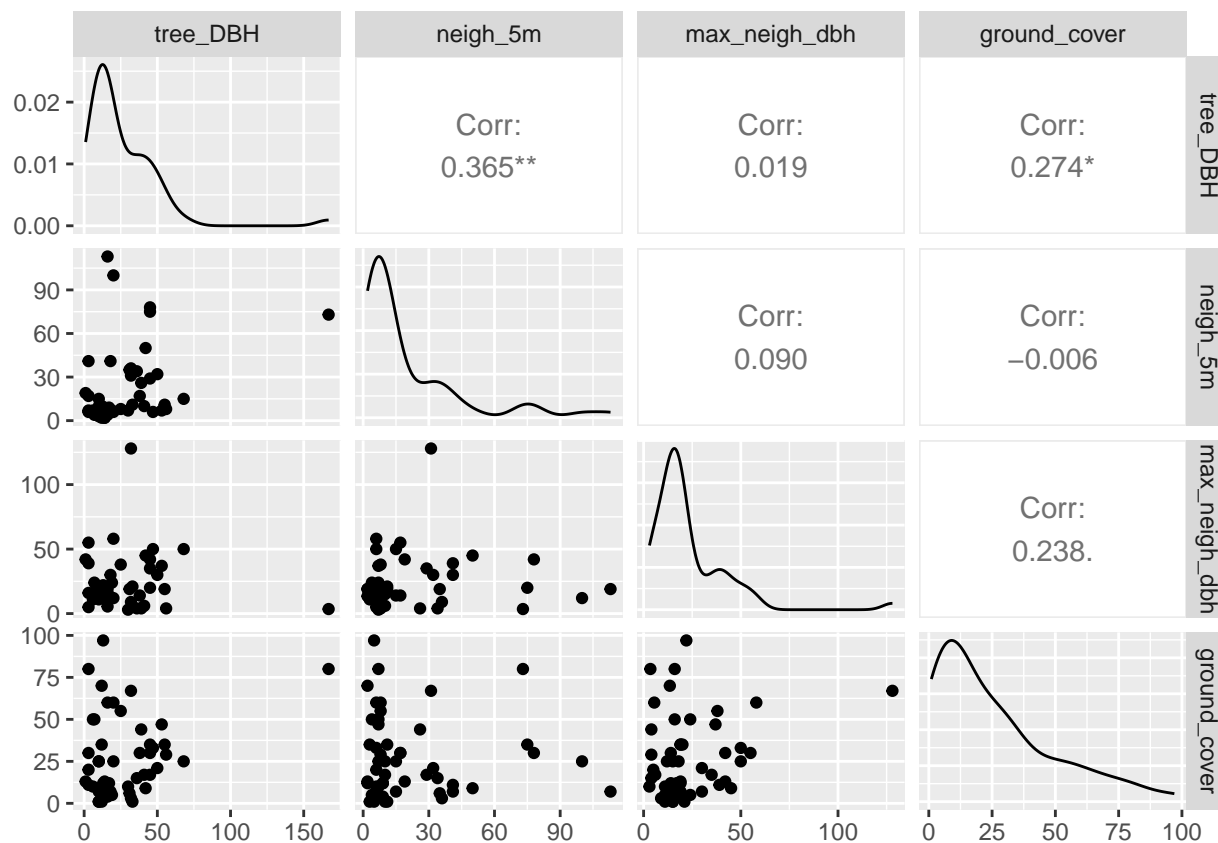
## Actual vs Predicted Tree DBH



```
# Load the GGally package for ggpairs  
library(GGally)
```

```
## Registered S3 method overwritten by 'GGally':  
##   method from  
##   +.gg      ggplot2
```

```
# Create a scatterplot matrix  
ggpairs(riseholme_data_24_1_[c("tree_DBH", "neigh_5m", "max_neigh_dbh", "ground_cover")])
```



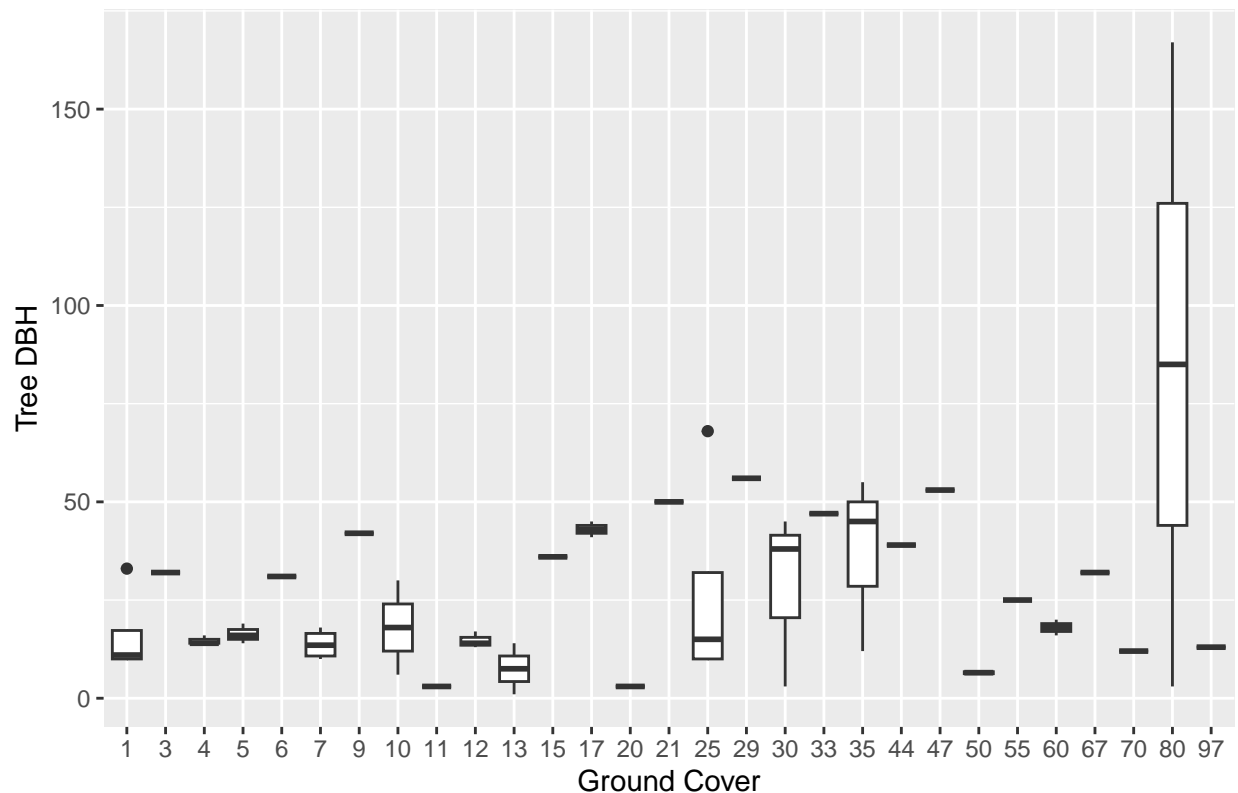
```
# Load the ggplot2 package
```

```
library(ggplot2)
```

```
# Create a boxplot of tree_DBH by ground_cover
```

```
ggplot(riseholme_data_24_1_, aes(x = as.factor(ground_cover), y = tree_DBH)) +  
  geom_boxplot() +  
  xlab("Ground Cover") +  
  ylab("Tree DBH") +  
  ggtitle("Distribution of Tree DBH by Ground Cover")
```

Distribution of Tree DBH by Ground Cover



```
# Create a density plot of tree_DBH by ground_cover
ggplot(riseholme_data_24_1, aes(x = tree_DBH, fill = as.factor(ground_cover))) +
  geom_density(alpha = 0.5) +
  xlab("Tree DBH") +
  ylab("Density") +
  ggtitle("Density Plot of Tree DBH by Ground Cover")
```

```
## Warning: Groups with fewer than two data points have been dropped.
## Groups with fewer than two data points have been dropped.
## Groups with fewer than two data points have been dropped.
## Groups with fewer than two data points have been dropped.
## Groups with fewer than two data points have been dropped.
## Groups with fewer than two data points have been dropped.
## Groups with fewer than two data points have been dropped.
## Groups with fewer than two data points have been dropped.
## Groups with fewer than two data points have been dropped.
## Groups with fewer than two data points have been dropped.
## Groups with fewer than two data points have been dropped.
## Groups with fewer than two data points have been dropped.
## Groups with fewer than two data points have been dropped.
## Groups with fewer than two data points have been dropped.
```

```
## Warning in max(ids, na.rm = TRUE): no non-missing arguments to max; returning
## -Inf
```

```
## Warning in max(ids, na.rm = TRUE): no non-missing arguments to max; returning
## -Inf
## Warning in max(ids, na.rm = TRUE): no non-missing arguments to max; returning
## -Inf
## Warning in max(ids, na.rm = TRUE): no non-missing arguments to max; returning
## -Inf
## Warning in max(ids, na.rm = TRUE): no non-missing arguments to max; returning
## -Inf
## Warning in max(ids, na.rm = TRUE): no non-missing arguments to max; returning
## -Inf
## Warning in max(ids, na.rm = TRUE): no non-missing arguments to max; returning
## -Inf
## Warning in max(ids, na.rm = TRUE): no non-missing arguments to max; returning
## -Inf
## Warning in max(ids, na.rm = TRUE): no non-missing arguments to max; returning
## -Inf
## Warning in max(ids, na.rm = TRUE): no non-missing arguments to max; returning
## -Inf
## Warning in max(ids, na.rm = TRUE): no non-missing arguments to max; returning
## -Inf
## Warning in max(ids, na.rm = TRUE): no non-missing arguments to max; returning
## -Inf
## Warning in max(ids, na.rm = TRUE): no non-missing arguments to max; returning
## -Inf
## Warning in max(ids, na.rm = TRUE): no non-missing arguments to max; returning
## -Inf
```

Density Plot of Tree DBH by Ground Cover

