Data importation

- 1. Endpoint dataframe
 - A. Exploration of data

Exploration tables using the rstatix, janitor and skimr packages

Data visualization

B. Normality hypothesis and outlier detection

Boxplots after outlier detection

Violin and sina plots after outlier detection

Exploration statistics for the variables after outlier detection

2. Exploration of the timeseries data

Number of data observations per day for the traits of the timeseries datasets

- A. Exploration of the timeseries dataframe
- B. Exploration of the S_timeseries dataframe
- C. Exploration of the T_timeseries dataframe

UCL Data Analysis

Elise

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Set the right working directory.

```
setwd("C:/Users/elise/Documents/Mémoire/Main/Data/Templates/UCL")
```

Data importation

Import the data sets extracted from the Data Preparation R Markdown.

```
list.files()
```

```
plant_info <- read.table("plant_info.txt", header = TRUE, sep = "\t")
endpoint <- read.table("endpoint.txt", header = TRUE, sep = "\t")</pre>
```

Convert the columns to factor and date formats.

```
# plant_info
plant_info <- lapply(plant_info, factor)

# endpoint
matching_cols <- intersect(names(endpoint), names(plant_info))
endpoint[, matching_cols] <- lapply(endpoint[, matching_cols], factor)
endpoint$Date <- date(endpoint$Date)
endpoint$Timestamp <- NA

# timeseries
# No data

# S_timeseries
# No data

# T_timeseries
# No data</pre>
```

Collect the variables of every data template and print the names of the variables. This serves as a double check.

```
platform <- "UCL"

# endpoint

df <- endpoint[,colSums(is.na(endpoint))<nrow(endpoint)]

genotype_index <- which(colnames(df) == "Genotype")

variables <- colnames(df[, c(3:(genotype_index - 1))]) # We remove the 3 first columns
that are "Unit.ID" and "Date" etc

print(paste(platform, ": The variables for endpoint are", paste(variables, collapse =
    ", "), sep = " "))</pre>
```

```
## [1] "UCL : The variables for endpoint are DW_shoot_g, FW_shoot_g, DW_root_g, FW_root_g"
```

Remove unknown genotypes

```
endpoint <- endpoint %>%
filter(Genotype != "Local")
```

Add a column Plant_type with three levels, H L and T. This variable is useful to test for heterosis effects.

```
endpoint$Plant_type <- substr(endpoint$Genotype, nchar(as.character(endpoint$Genotyp
e)), nchar(as.character(endpoint$Genotype)))</pre>
```

1. Endpoint dataframe

A. Exploration of data

Exploration tables using the rstatix, janitor and skimr packages

```
endpoint %>%
count(Genotype)
```

```
##
     Genotype n
## 1 EPPN1_H 29
## 2 EPPN1_L 23
## 3 EPPN10 H 25
## 4 EPPN10 L 10
## 5 EPPN11_H 27
## 6 EPPN11_L 21
## 7 EPPN12_H 25
## 8 EPPN12_L 14
## 9 EPPN13_H 15
## 10 EPPN13_L 6
## 11 EPPN14 H 15
## 12 EPPN14_L 6
## 13 EPPN15_H 25
## 14 EPPN15_L 18
## 15 EPPN2_H 24
## 16 EPPN2_L 17
## 17 EPPN20_T 27
## 18 EPPN3_H 18
## 19 EPPN3_L 19
## 20 EPPN4_H 20
## 21 EPPN4_L 20
## 22 EPPN5 H 16
## 23 EPPN5_L 8
## 24 EPPN6_H 24
## 25 EPPN6 L 17
## 26 EPPN7_H 7
## 27 EPPN7_L 16
## 28 EPPN8_H 21
## 29 EPPN8 L 18
## 30 EPPN9 H 25
## 31 EPPN9_L 22
```

```
endpoint %>%
  tabyl(Genotype, Column) %>%
  adorn_totals("row") %>%
  adorn_percentages("row") %>%
  adorn_pct_formatting() %>%
  adorn_ns() %>%
  adorn_ns() %>%
```

```
5
##
    Genotype/Column
                                1
                                            2
                                                                     4
                                                         3
             EPPN1_H 10.3%
                              (3) 10.3%
                                          (3) 10.3%
##
                                                      (3)
                                                            6.9%
                                                                   (2) 10.3%
                                                                               (3)
##
             EPPN1 L
                       8.7%
                              (2)
                                   8.7%
                                          (2)
                                                8.7%
                                                       (2)
                                                            8.7%
                                                                   (2) 13.0%
                                                                               (3)
##
            EPPN10 H 8.0%
                              (2)
                                   8.0%
                                          (2)
                                                8.0%
                                                       (2)
                                                            8.0%
                                                                   (2) 12.0%
                                                                               (3)
            EPPN10 L 10.0%
                              (1) 20.0%
                                          (2) 20.0%
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                                                            0.0%
                                                                   (0)
                                                                        0.0%
                                                                               (0)
##
##
            EPPN11_H 11.1%
                              (3) 11.1%
                                          (3)
                                                7.4%
                                                       (2)
                                                            7.4%
                                                                   (2) 11.1%
                                                                               (3)
##
            EPPN11_L 14.3%
                              (3)
                                   4.8%
                                          (1)
                                                9.5%
                                                       (2)
                                                            0.0%
                                                                   (0) 14.3%
                                                                               (3)
                                                       (2)
##
            EPPN12_H 12.0%
                              (3) 12.0%
                                          (3)
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                                                            8.0%
                                                                   (2)
                                                                        8.0%
                                                                               (2)
##
            EPPN12 L 7.1%
                              (1)
                                   7.1%
                                          (1) 14.3%
                                                       (2)
                                                            7.1%
                                                                   (1) 21.4%
                                                                               (3)
##
            EPPN13_H 13.3%
                              (2) 13.3%
                                          (2) 13.3%
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                                                            6.7%
                                                                   (1) 20.0%
                                                                               (3)
            EPPN13 L 16.7%
                              (1) 16.7%
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##
                                                0.0%
                                                                               (1)
            EPPN14 H 6.7%
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                                          (2) 20.0%
                                                       (3) 13.3%
                                                                   (2)
##
                                                                        6.7%
                                                                               (1)
##
            EPPN14_L 33.3%
                              (2) 16.7%
                                          (1) 16.7%
                                                      (1) 16.7%
                                                                   (1) 16.7%
                                                                               (1)
            EPPN15_H 8.0%
                              (2) 12.0%
                                          (3) 12.0%
                                                       (3)
                                                            8.0%
                                                                   (2) 12.0%
                                                                               (3)
##
            EPPN15_L 11.1%
##
                              (2)
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                                          (1)
             EPPN2_H 12.5%
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             EPPN2_L
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##
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             EPPN3 H
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##
##
             EPPN3_L 10.5%
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##
             EPPN4_H
                       5.0%
                                          (3) 10.0%
                                                       (2) 10.0%
                                                                   (2) 15.0%
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##
             EPPN4_L
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                                   5.0%
                                          (1) 10.0%
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##
             EPPN5 H 6.2%
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                                   6.2%
                                          (1) 12.5%
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                                                                   (2) 18.8%
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##
             EPPN5_L 12.5%
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##
             EPPN6_H 12.5%
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##
             EPPN7 H 28.6%
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             EPPN7 L 12.5%
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                                                                               (0)
##
             EPPN8_H 9.5%
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                                   9.5%
                                          (2)
                                                9.5%
                                                      (2)
                                                            4.8%
                                                                   (1)
                                                                        9.5%
                                                                               (2)
##
             EPPN8 L 11.1%
                              (2) 16.7%
                                          (3) 16.7%
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##
             EPPN9 H
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##
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                                          (2) 13.6%
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##
               Local
                                                                               (0)
##
               Total
                       9.9%
                            (57)
                                   9.2%
                                         (53) 10.2%
                                                     (59)
                                                            9.5%
                                                                 (55) 11.8% (68)
##
              6
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                                       8
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##
    10.3%
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##
            (3)
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                                     (2) 14.3%
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                                                      7.1%
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                                     (1)
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    16.7%
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                                                             (0)
##
     6.7%
            (1)
                 6.7%
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                              0.0%
                                     (0) 13.3%
                                                 (2) 13.3%
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##
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##
    12.0%
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                 4.0%
                        (1) 12.0%
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                                          8.0%
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    11.1%
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                                     (2) 16.7%
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##
    17.6%
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##
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##
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                        (1) 15.8%
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```

```
##
   15.0% (3) 10.0% (2) 10.0% (2) 15.0% (3) 10.0%
                                                  (2)
##
   12.5% (2) 12.5% (2) 12.5% (2) 0.0% (0) 6.2%
                                                  (1)
   12.5% (1) 25.0% (2) 0.0% (0) 12.5% (1) 12.5%
##
                                                  (1)
    8.3% (2) 4.2% (1) 12.5% (3) 8.3% (2) 12.5%
##
                                                 (3)
##
    5.9% (1) 5.9%
                   (1) 11.8% (2) 11.8% (2) 11.8%
                                                  (2)
##
    0.0% (0) 14.3%
                   (1) 14.3% (1) 0.0% (0) 14.3% (1)
##
    6.2% (1) 12.5%
                   (2) 12.5% (2) 12.5% (2) 12.5% (2)
##
   14.3% (3) 9.5%
                   (2) 14.3% (3) 9.5%
                                       (2) 9.5% (2)
##
   11.1%
          (2) 11.1%
                   (2) 0.0%
                             (0) 5.6%
                                        (1)
                                            5.6%
                                                  (1)
##
   12.0% (3) 8.0%
                   (2) 12.0% (3) 8.0%
                                       (2) 8.0%
                                                 (2)
##
   13.6% (3) 9.1%
                   (2) 4.5% (1) 13.6% (3)
                                            4.5% (1)
##
         (0)
                   (0)
                           - (0)
                                     - (0)
                                               - (0)
   10.0% (58) 9.0% (52) 9.9% (57) 10.0% (58) 10.6% (61)
##
```

```
endpoint %>%
  tabyl(Genotype, Row) %>%
  adorn_totals("row") %>%
  adorn_percentages("row") %>%
  adorn_pct_formatting() %>%
  adorn_ns() %>%
  adorn_ns() %>%
```

```
##
    Genotype/Row
                         1
                                   2
                                             3
                                                                5
                                                                          6
                                                                                   7
         EPPN1_H 3.4% (1)
                            0.0% (0) 0.0% (0)
                                                3.4% (1) 0.0% (0) 3.4% (1) 0.0% (0)
##
         EPPN1 L 0.0% (0)
                            0.0% (0) 0.0% (0)
                                                0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
##
        EPPN10 H 0.0% (0)
                            0.0% (0) 4.0% (1)
                                                4.0% (1) 4.0% (1) 0.0% (0) 0.0% (0)
##
        EPPN10 L 0.0% (0)
                            0.0% (0) 0.0% (0)
                                                0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
##
##
        EPPN11_H 0.0% (0)
                            3.7% (1) 0.0% (0)
                                                0.0% (0) 0.0% (0) 3.7% (1) 3.7% (1)
##
        EPPN11_L 0.0% (0)
                            0.0% (0) 0.0% (0)
                                                0.0% (0) 4.8% (1) 0.0% (0) 0.0% (0)
        EPPN12 H 0.0% (0)
                            0.0% (0) 4.0% (1)
                                                4.0% (1) 0.0% (0) 0.0% (0) 0.0% (0)
##
        EPPN12_L 0.0% (0)
                            0.0% (0) 0.0% (0)
                                                0.0% (0) 7.1% (1) 0.0% (0) 0.0% (0)
##
##
        EPPN13_H 0.0% (0)
                            0.0% (0) 0.0% (0)
                                                0.0% (0) 6.7% (1) 0.0% (0) 6.7% (1)
        EPPN13 L 0.0% (0)
                            0.0% (0) 0.0% (0)
                                                0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
##
        EPPN14 H 6.7% (1)
                            0.0% (0) 0.0% (0)
                                                0.0% (0) 0.0% (0) 6.7% (1) 0.0% (0)
##
        EPPN14_L 0.0% (0) 16.7% (1) 0.0% (0)
                                                0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
##
        EPPN15_H 4.0% (1)
                            0.0% (0) 0.0% (0)
                                                0.0% (0) 4.0% (1) 0.0% (0) 0.0% (0)
##
        EPPN15_L 0.0% (0)
                            0.0% (0) 0.0% (0)
                                                0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
##
         EPPN2_H 4.2% (1)
                            4.2% (1) 4.2% (1)
                                                0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
##
         EPPN2_L 0.0% (0)
                            0.0% (0) 5.9% (1)
                                                0.0% (0) 5.9% (1) 0.0% (0) 0.0% (0)
##
        EPPN20_T 0.0% (0)
                                                0.0% (0) 0.0% (0) 0.0% (0) 3.7% (1)
##
                            0.0% (0) 0.0% (0)
         EPPN3 H 0.0% (0)
                                                0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
                            0.0% (0) 0.0% (0)
##
##
         EPPN3_L 0.0% (0) 10.5% (2) 0.0% (0)
                                                0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
##
         EPPN4_H 0.0% (0)
                            5.0% (1) 0.0% (0)
                                                0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
##
         EPPN4_L 5.0% (1)
                            0.0% (0) 0.0% (0)
                                                0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
##
         EPPN5 H 0.0% (0)
                            0.0% (0) 0.0% (0)
                                                0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
         EPPN5 L 0.0% (0)
                            0.0% (0) 0.0% (0)
                                                0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
##
         EPPN6 H 4.2% (1)
                            4.2% (1) 0.0% (0)
                                                0.0% (0) 0.0% (0) 0.0% (0) 4.2% (1)
##
         EPPN6 L 0.0% (0)
                                                5.9% (1) 0.0% (0) 0.0% (0) 0.0% (0)
##
                            0.0% (0) 0.0% (0)
##
         EPPN7 H 0.0% (0)
                            0.0% (0) 0.0% (0) 14.3% (1) 0.0% (0) 0.0% (0) 0.0% (0)
         EPPN7 L 0.0% (0)
                            6.2% (1) 0.0% (0)
                                                0.0% (0) 0.0% (0) 6.2% (1) 0.0% (0)
##
         EPPN8_H 0.0% (0)
                            0.0% (0) 4.8% (1)
                                                4.8% (1) 0.0% (0) 0.0% (0) 0.0% (0)
##
         EPPN8 L 0.0% (0)
                            0.0% (0) 5.6% (1)
                                                5.6% (1) 5.6% (1) 0.0% (0) 5.6% (1)
##
                                                0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
##
         EPPN9_H 0.0% (0)
                            0.0% (0) 4.0% (1)
         EPPN9_L 0.0% (0)
##
                            4.5% (1) 0.0% (0)
                                                0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
           Local
                     - (0)
                               - (0)
                                         - (0)
                                                   - (0)
                                                            - (0)
                                                                      - (0)
##
                                                                               - (0)
##
           Total 1.0% (6)
                            1.6% (9) 1.2% (7)
                                                1.2% (7) 1.2% (7) 0.7% (4) 0.9% (5)
##
                                10
                                          11
                                                   12
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                         1.7% (10) 1.2% (7) 0.9% (5) 1.4% (8) 1.0% (6)
##
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##
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##
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##
##
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##
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##
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##
##
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##
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##
##
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##
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                                                      1.4% (8)
                                                                1.6% (9) 0.9% (5)
##
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##
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##
    3.4% (1) 0.0% (0)
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##
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##
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```

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##
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##
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##
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##
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##
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##
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##
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##
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##
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##
##
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    0.0% (0) 0.0% (0) 11.1% (2)
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##
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##
##
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    0.9% (5) 1.4% (8) 0.7% (4)
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                                                     1.0% (6) 0.9% (5) 1.2% (7)
##
##
          32
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                                                           37
                                                                     38
##
    3.4% (1)
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##
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##
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##
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##
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##
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                                                               7.1% (1) 7.1% (1)
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##
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                                                             0.0% (0) 4.0% (1)
##
                       0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
##
    0.0% (0) 0.0% (0)
                                                             0.0% (0) 0.0% (0)
    0.0% (0) 0.0% (0)
                       0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
                                                             0.0% (0) 6.7% (1)
##
    0.0% (0) 0.0% (0)
                       0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0) 16.7% (1) 0.0% (0)
##
##
    0.0% (0) 0.0% (0)
                       0.0% (0) 0.0% (0) 6.7% (1) 0.0% (0)
                                                             0.0% (0) 0.0% (0)
                       0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
##
    0.0% (0) 0.0% (0)
                                                             0.0% (0) 0.0% (0)
##
    0.0% (0) 0.0% (0)
                       0.0% (0) 0.0% (0) 4.0% (1) 0.0% (0)
                                                             0.0% (0) 0.0% (0)
    0.0% (0) 0.0% (0)
                       0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
                                                             0.0% (0) 0.0% (0)
##
##
    0.0% (0) 0.0% (0)
                       4.2% (1) 0.0% (0) 0.0% (0) 0.0% (0)
                                                             0.0% (0) 0.0% (0)
   0.0% (0) 0.0% (0)
                       0.0% (0) 5.9% (1) 0.0% (0) 0.0% (0)
                                                             0.0% (0) 0.0% (0)
```

```
##
   3.7% (1) 0.0% (0) 3.7% (1) 3.7% (1) 0.0% (0) 3.7% (1) 0.0% (0) 0.0% (0)
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0) 5.6% (1) 0.0% (0)
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
                                                           0.0% (0) 0.0% (0)
##
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0) 5.0% (1) 0.0% (0)
                                                           0.0% (0) 0.0% (0)
                      0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
##
   0.0% (0) 5.0% (1)
                                                           0.0% (0) 5.0% (1)
   6.2% (1) 0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
                                                           0.0% (0) 0.0% (0)
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
                                                           0.0% (0) 0.0% (0)
##
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
                                                           0.0% (0) 8.3% (2)
##
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0) 5.9% (1) 0.0% (0)
                                                           5.9% (1) 0.0% (0)
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
                                                           0.0% (0) 0.0% (0)
   0.0% (0) 6.2% (1)
                      0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
                                                           0.0% (0) 0.0% (0)
##
   0.0% (0) 0.0% (0)
                      4.8% (1) 0.0% (0) 0.0% (0) 0.0% (0)
                                                           0.0% (0) 0.0% (0)
##
    0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
                                                           0.0% (0) 5.6% (1)
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0) 4.0% (1)
                                                           0.0% (0) 0.0% (0)
    0.0% (0) 4.5% (1) 0.0% (0) 0.0% (0) 4.5% (1) 0.0% (0)
                                                           0.0% (0) 0.0% (0)
##
##
               - (0)
                         - (0)
                                  - (0)
                                          - (0)
                                                   - (0)
                                                              - (0)
      - (0)
                                                                       - (0)
   0.7% (4) 0.9% (5) 1.0% (6) 0.7% (4) 1.2% (7) 0.3% (2) 0.7% (4) 1.4% (8)
##
                                     99
         96
                  97
                            98
##
   0.0% (0) 0.0% (0) 0.0% (0) 3.4% (1)
##
##
   0.0% (0) 0.0% (0) 0.0% (0) 4.3% (1)
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
##
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
##
##
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
   4.8% (1) 0.0% (0) 0.0% (0) 0.0% (0)
##
   0.0% (0) 0.0% (0) 8.0% (2) 0.0% (0)
##
   0.0% (0) 0.0% (0) 7.1% (1) 0.0% (0)
##
##
   0.0% (0) 0.0% (0) 0.0% (0) 6.7% (1)
##
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
##
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
##
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
   4.2% (1) 0.0% (0) 0.0% (0) 0.0% (0)
##
   5.9% (1) 5.9% (1) 0.0% (0) 0.0% (0)
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
    5.6% (1) 0.0% (0) 0.0% (0) 0.0% (0)
##
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
##
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
   0.0% (0) 5.0% (1) 0.0% (0) 0.0% (0)
##
    6.2% (1) 0.0% (0) 0.0% (0) 6.2% (1)
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
##
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
   0.0% (0) 5.9% (1) 0.0% (0) 0.0% (0)
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
##
##
   0.0% (0) 6.2% (1) 0.0% (0) 0.0% (0)
   0.0% (0) 0.0% (0) 0.0% (0) 4.8% (1)
##
##
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
   0.0% (0) 0.0% (0) 0.0% (0) 0.0% (0)
##
   0.0% (0) 0.0% (0) 0.0% (0) 4.5% (1)
##
      - (0)
              - (0) - (0)
##
                                - (0)
##
   0.9% (5) 0.7% (4) 0.5% (3) 1.0% (6)
```

```
## Warning: Using an external vector in selections was deprecated in tidyselect 1.1.0.
## i Please use `all_of()` or `any_of()` instead.
## # Was:
## data %>% select(variables)
##
## # Now:
## data %>% select(all_of(variables))
##
## See <https://tidyselect.r-lib.org/reference/faq-external-vector.html>.
## This warning is displayed once every 8 hours.
## Call `lifecycle::last_lifecycle_warnings()` to see where this warning was
## generated.
```

```
## # A tibble: 4 × 10
   variable
##
                  min
                       max median
                                 iqr mean
   ##
                    0 0.538 0.238 0.122 0.24 0.095 0.004 0.008
## 1 DW_shoot_g 572
## 2 FW_shoot_g 574
                    0 8.63 3.48 2.07 3.52 1.60 0.067 0.131
                    0 0.275 0.102 0.056 0.104 0.048 0.002 0.004
## 3 DW_root_g
             577
## 4 FW_root_g
             576
                    0 7.29
                           2.52 1.78 2.57 1.33 0.055 0.109
```

```
skim(endpoint[variables])
```

Data summary

Name	endpoint[variables]
Number of rows	578
Number of columns	4
Column type frequency:	
numeric	4
Group variables	None

Variable type: numeric

skim_variable	n_missing	complete_rate	mean	sd	p0	p25	p50	p75	p100	hist
DW_shoot_g	6	0.99	0.24	0.09	0	0.18	0.24	0.30	0.54	_===_
FW_shoot_g	4	0.99	3.52	1.60	0	2.51	3.48	4.58	8.63	_==_
DW_root_g	1	1.00	0.10	0.05	0	0.07	0.10	0.13	0.28	_==_
FW_root_g	2	1.00	2.57	1.33	0	1.65	2.51	3.43	7.29	_===

Data visualization

Using several functions that are located in the functions.R script

Boxplots

```
create_boxplots(endpoint, variables, "Genotype")
```

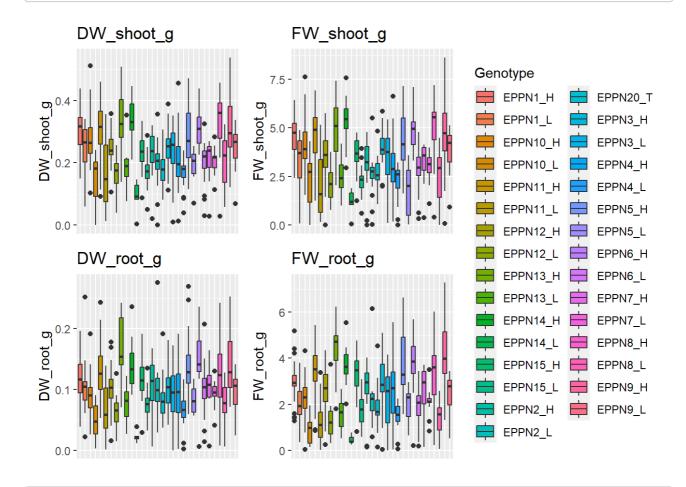
```
## Warning: `aes_string()` was deprecated in ggplot2 3.0.0.
## i Please use tidy evaluation idioms with `aes()`.
## i See also `vignette("ggplot2-in-packages")` for more information.
## This warning is displayed once every 8 hours.
## Call `lifecycle::last_lifecycle_warnings()` to see where this warning was
## generated.
```

Warning: Removed 6 rows containing non-finite values (`stat_boxplot()`).

Warning: Removed 4 rows containing non-finite values (`stat_boxplot()`).

Warning: Removed 1 rows containing non-finite values (`stat_boxplot()`).

Warning: Removed 2 rows containing non-finite values (`stat_boxplot()`).



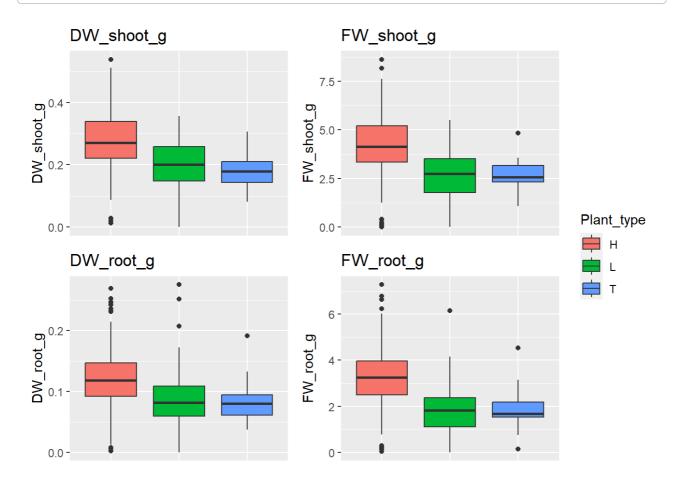
create_boxplots(endpoint, variables, "Plant_type")

Warning: Removed 6 rows containing non-finite values (`stat_boxplot()`).

Warning: Removed 4 rows containing non-finite values (`stat_boxplot()`).

```
## Warning: Removed 1 rows containing non-finite values (`stat_boxplot()`).
```

Warning: Removed 2 rows containing non-finite values (`stat_boxplot()`).



Correlation plots

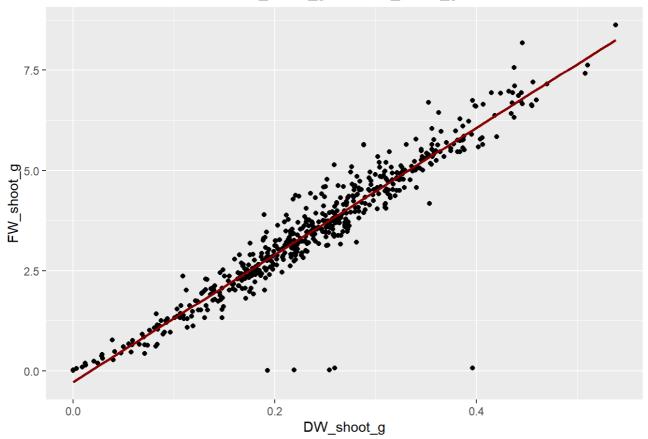
```
for (i in 1:(length(variables) - 1)) {
  for (j in (i + 1):length(variables)) {
    calculate_correlation_plot(endpoint, variables[i], variables[j])
  }
}
```

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

Warning: Removed 7 rows containing non-finite values (`stat_smooth()`).

```
## Warning: Removed 7 rows containing missing values (`geom_point()`).
```

Correlation Plot between DW_shoot_g and FW_shoot_g

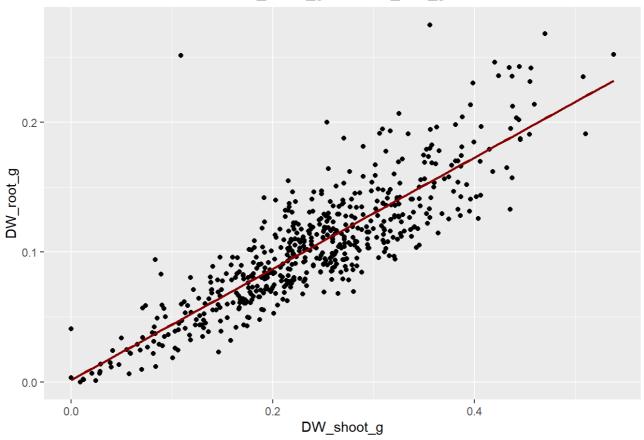


$geom_smooth()$ using formula = 'y ~ x'

Warning: Removed 7 rows containing non-finite values (`stat_smooth()`).

Warning: Removed 7 rows containing missing values (`geom_point()`).

Correlation Plot between DW_shoot_g and DW_root_g

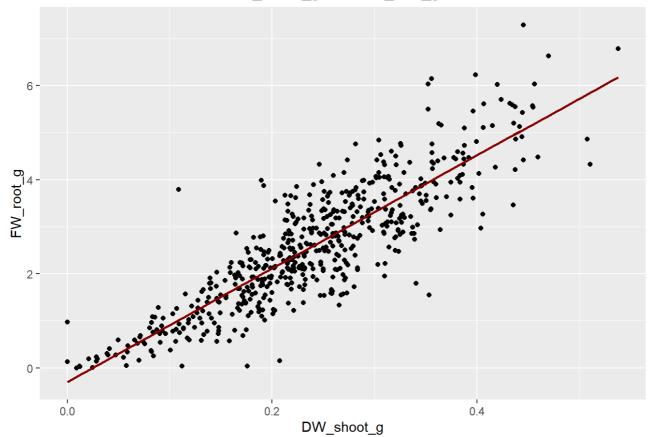


`geom_smooth()` using formula = 'y ~ x'

Warning: Removed 8 rows containing non-finite values (`stat_smooth()`).

Warning: Removed 8 rows containing missing values (`geom_point()`).

Correlation Plot between DW_shoot_g and FW_root_g

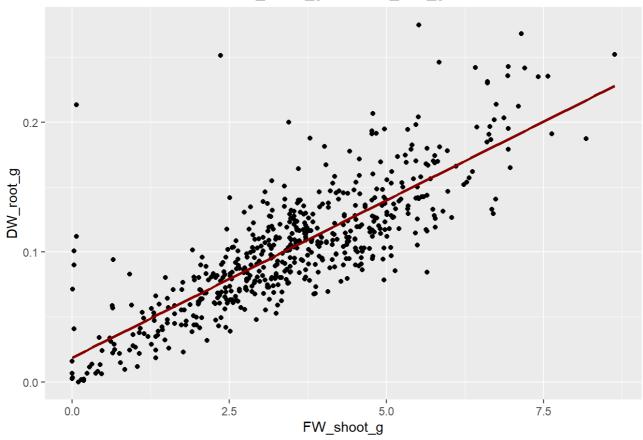


$geom_smooth()$ using formula = 'y ~ x'

Warning: Removed 5 rows containing non-finite values (`stat_smooth()`).

Warning: Removed 5 rows containing missing values (`geom_point()`).

Correlation Plot between FW_shoot_g and DW_root_g

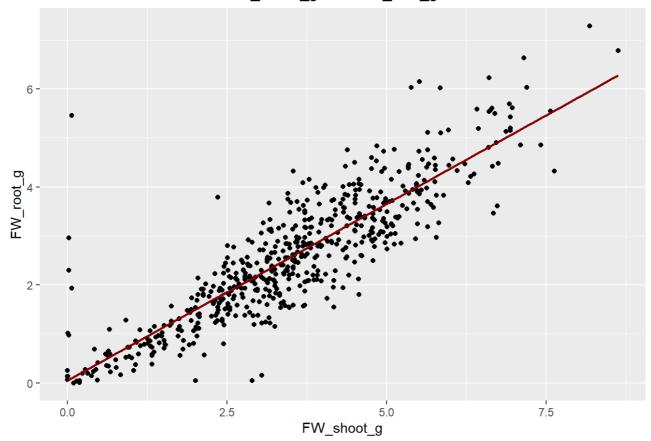


$geom_smooth()$ using formula = 'y ~ x'

Warning: Removed 6 rows containing non-finite values (`stat_smooth()`).

Warning: Removed 6 rows containing missing values (`geom_point()`).

Correlation Plot between FW_shoot_g and FW_root_g

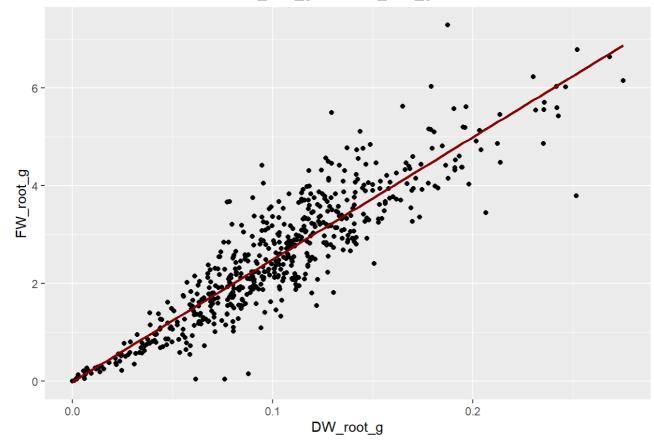


$geom_smooth()$ using formula = 'y ~ x'

Warning: Removed 2 rows containing non-finite values (`stat_smooth()`).

Warning: Removed 2 rows containing missing values (`geom_point()`).

Correlation Plot between DW_root_g and FW_root_g



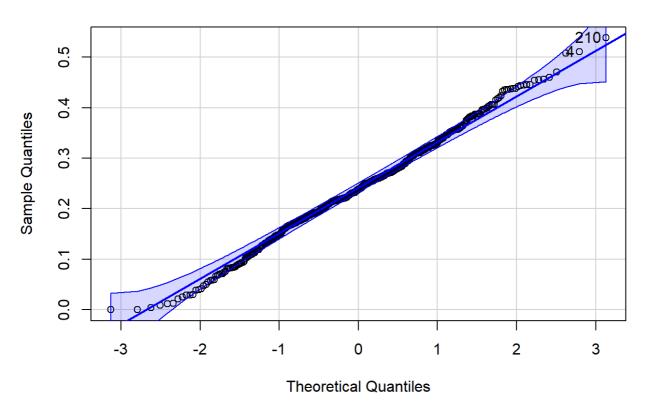
B. Normality hypothesis and outlier detection

Test for normality hypothesis and plot density histogram. The red curve is the normal distribution, the blue dotted curve is the data density curve.

```
normality_results <- normality_test_histogram(endpoint)</pre>
```

```
## Warning: Using `size` aesthetic for lines was deprecated in ggplot2 3.4.0.
## i Please use `linewidth` instead.
## This warning is displayed once every 8 hours.
## Call `lifecycle::last_lifecycle_warnings()` to see where this warning was
## generated.
```

QQ Plot of DW_shoot_g

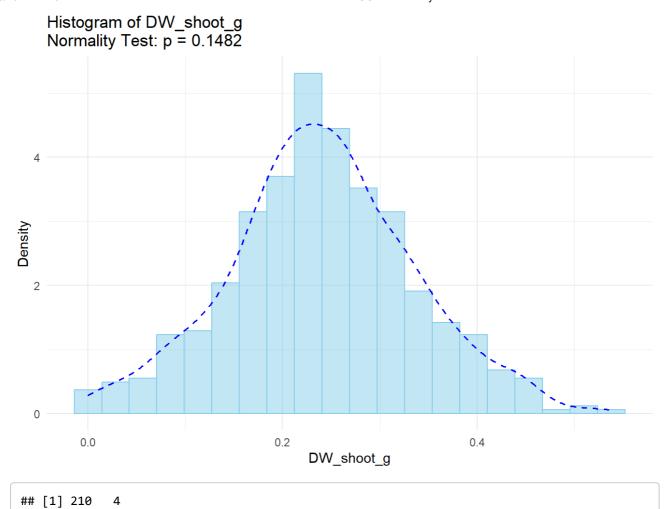


```
## Warning: The dot-dot notation (`..density..`) was deprecated in ggplot2 3.4.0.
## i Please use `after_stat(density)` instead.
## This warning is displayed once every 8 hours.
## Call `lifecycle::last_lifecycle_warnings()` to see where this warning was
## generated.
```

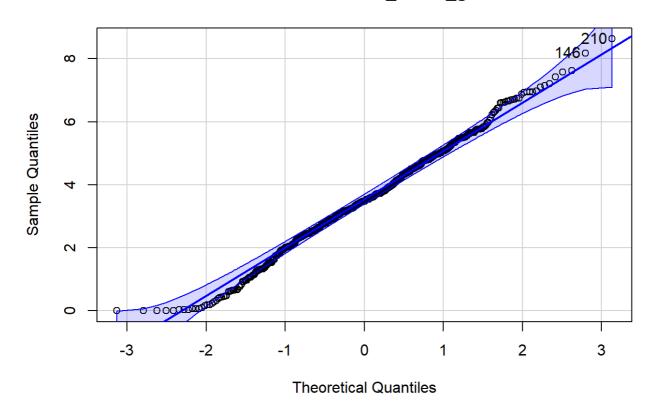
```
## Warning: Removed 6 rows containing non-finite values (`stat_bin()`).
```

```
## Warning: Removed 6 rows containing non-finite values (`stat_density()`).
```

Warning: Removed 101 rows containing missing values (`geom_function()`).



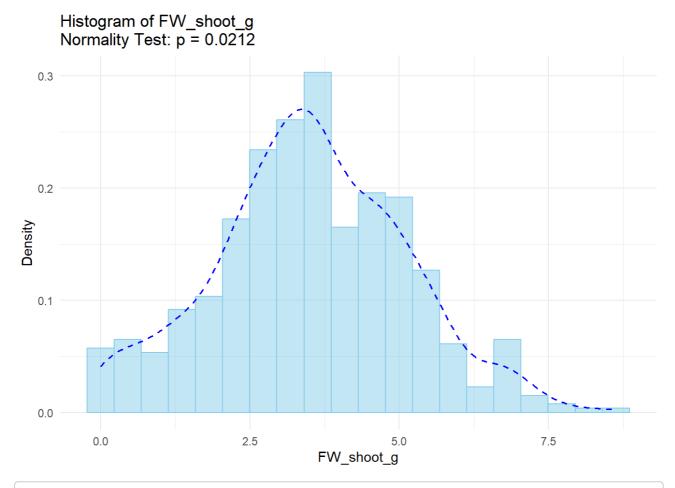
QQ Plot of FW_shoot_g



Warning: Removed 4 rows containing non-finite values (`stat_bin()`).

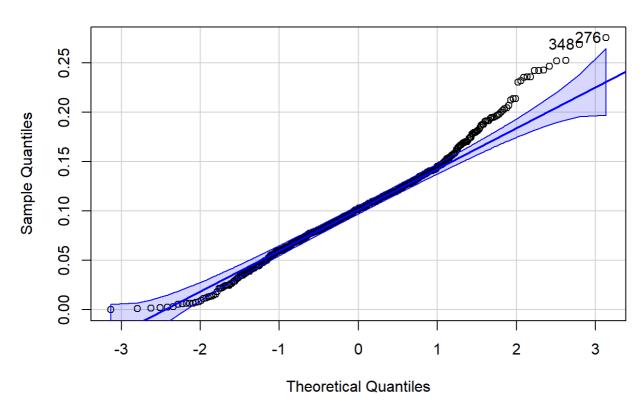
Warning: Removed 4 rows containing non-finite values (`stat_density()`).

Warning: Removed 101 rows containing missing values (`geom_function()`).



[1] 210 146

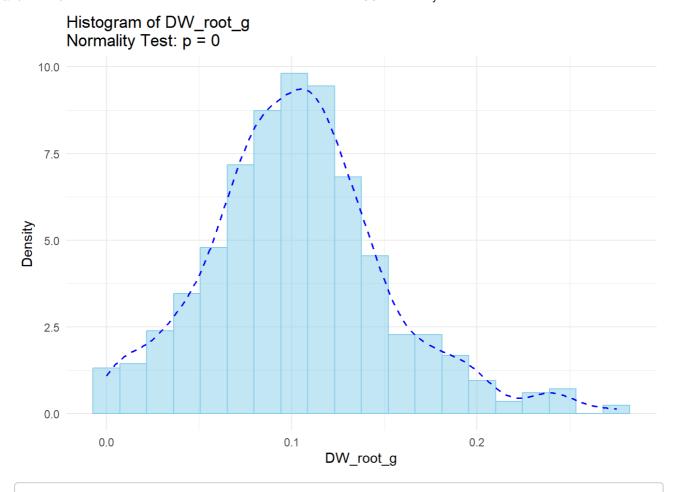
QQ Plot of DW_root_g

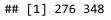


Warning: Removed 1 rows containing non-finite values (`stat_bin()`).

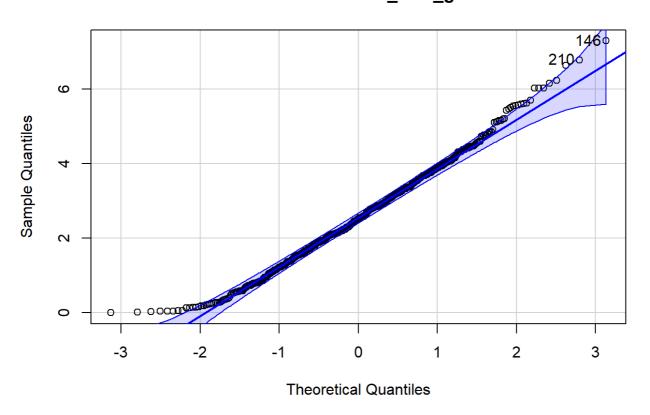
Warning: Removed 1 rows containing non-finite values (`stat_density()`).

Warning: Removed 101 rows containing missing values (`geom_function()`).





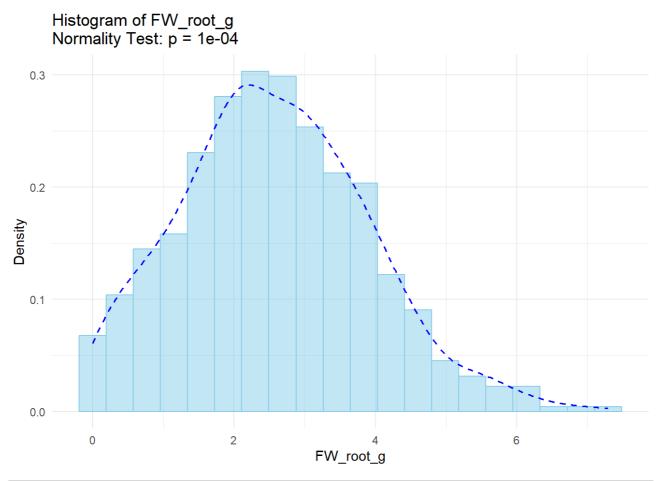




Warning: Removed 2 rows containing non-finite values (`stat_bin()`).

Warning: Removed 2 rows containing non-finite values (`stat_density()`).

Warning: Removed 101 rows containing missing values (`geom_function()`).



```
## [1] 146 210
```

Remove the outliers, replacing them with NULL values and normality visual verification.

The function detect_replace_ouliers_by_genotype checks for outlying values, using the Tukey method.

Then run the function on all variables of the dataset.

```
endpoint_clean <- endpoint
# Run the function on the dataset for all the variables
endpoint_clean <- detect_replace_outliers_by_genotype(endpoint_clean)</pre>
```

Boxplots after outlier detection

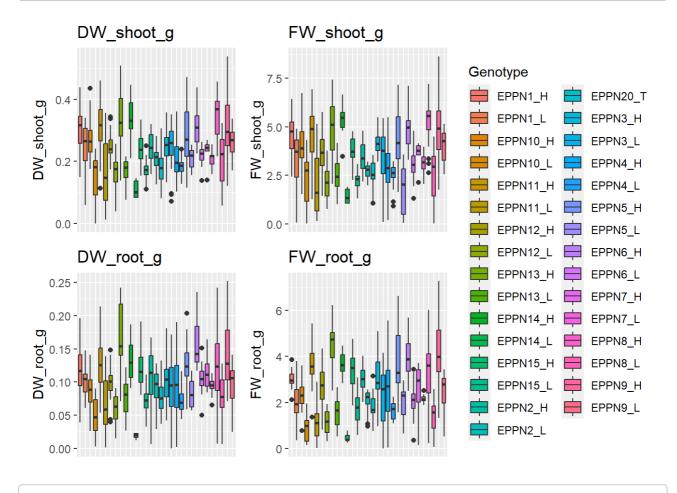
```
create_boxplots(endpoint_clean, variables, "Genotype")

## Warning: Removed 38 rows containing non-finite values (`stat_boxplot()`).

## Warning: Removed 34 rows containing non-finite values (`stat_boxplot()`).
```

Warning: Removed 26 rows containing non-finite values (`stat_boxplot()`).

Warning: Removed 34 rows containing non-finite values (`stat_boxplot()`).

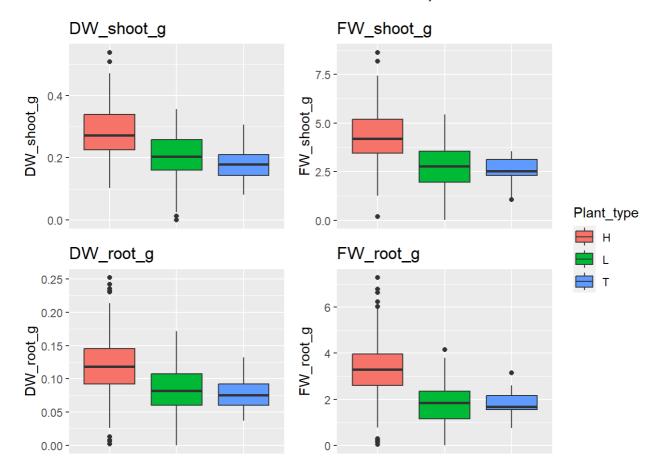


create_boxplots(endpoint_clean, variables, "Plant_type")

Warning: Removed 38 rows containing non-finite values (`stat_boxplot()`).
Removed 34 rows containing non-finite values (`stat_boxplot()`).

Warning: Removed 26 rows containing non-finite values (`stat_boxplot()`).

Warning: Removed 34 rows containing non-finite values (`stat_boxplot()`).



Violin and sina plots after outlier detection

```
## Warning: The `size` argument of `element_line()` is deprecated as of ggplot2 3.4.0.
## i Please use the `linewidth` argument instead.
## This warning is displayed once every 8 hours.
## Call `lifecycle::last_lifecycle_warnings()` to see where this warning was
## generated.

## Warning: Removed 38 rows containing non-finite values (`stat_ydensity()`).

## Warning: Removed 34 rows containing non-finite values (`stat_sina()`).

## Warning: Removed 34 rows containing non-finite values (`stat_sina()`).

## Warning: Removed 36 rows containing non-finite values (`stat_sina()`).

## Warning: Removed 26 rows containing non-finite values (`stat_ydensity()`).

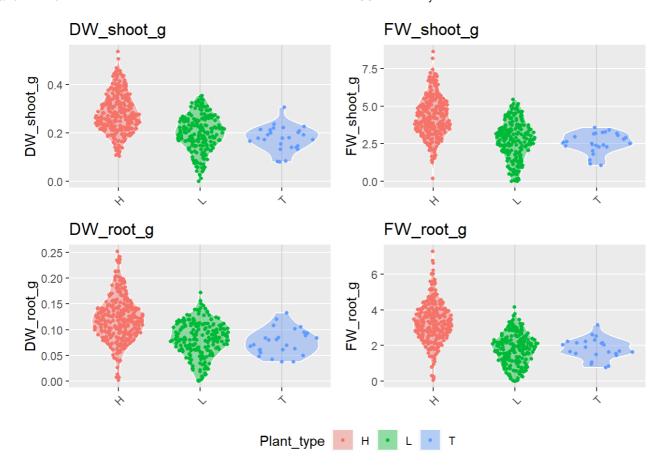
## Warning: Removed 26 rows containing non-finite values (`stat_ydensity()`).

## Warning: Removed 26 rows containing non-finite values (`stat_sina()`).

## Warning: Removed 34 rows containing non-finite values (`stat_sina()`).
```

Warning: Removed 34 rows containing non-finite values (`stat_sina()`).

```
DW shoot_g
       DW shoot g
                                                    FW shoot g
       DW root g
                                                    FW root_g
                                              FW_root_g
DW root
                         EPPN1_H
                                      EPPN12_L
                                                    EPPN2_H
                                                                 EPPN5_H
                                                                              EPPN8_L
                                      EPPN13_H
                         EPPN1_L
                                                    EPPN2_L
                                                                 EPPN5 L
                                                                              EPPN9 H
                         EPPN10_H
                                      EPPN13_L
                                                    EPPN20_T
                                                                 EPPN6_H
                                                                             EPPN9_L
                         EPPN10_L
           Genotype
                                      EPPN14_H
                                                    EPPN3_H
                                                                 EPPN6 L
                         EPPN11_H
                                      EPPN14_L
                                                    EPPN3_L
                                                                 EPPN7_H
                                      EPPN15 H
                                                    EPPN4 H
                                                                 EPPN7 L
                         EPPN11_L
                                                    EPPN4_L
                         EPPN12_H
                                      EPPN15_L
                                                                 EPPN8_H
create violin plots(endpoint clean, variables, "Plant type")
## Warning: Removed 38 rows containing non-finite values (`stat_ydensity()`).
## Warning: Removed 38 rows containing non-finite values (`stat sina()`).
## Warning: Removed 34 rows containing non-finite values (`stat_ydensity()`).
## Warning: Removed 34 rows containing non-finite values (`stat_sina()`).
## Warning: Removed 26 rows containing non-finite values (`stat ydensity()`).
## Warning: Removed 26 rows containing non-finite values (`stat_sina()`).
## Warning: Removed 34 rows containing non-finite values (`stat ydensity()`).
## Warning: Removed 34 rows containing non-finite values (`stat sina()`).
```



Exploration statistics for the variables after outlier detection

skim(endpoint_clean[variables])

Data summary

Name	endpoint_clean[variables]
Number of rows	578
Number of columns	4
Column type frequency:	
numeric	4
Group variables	None

Variable type: numeric

skim_variable	n_missing	complete_rate	mean	sd	p0	p25	p50	p75	p100	hist	
DW_shoot_g	38	0.93	0.25	0.09	0.00	0.19	0.24	0.30	0.54		
FW_shoot_g	34	0.94	3.59	1.51	0.01	2.61	3.53	4.60	8.63	_==_	_
DW_root_g	26	0.96	0.10	0.04	0.00	0.07	0.10	0.13	0.25	_==_	_

skim_variable n_missing complete_rate mean sd p0 p25 p50 p75 p100 hist

FW_root_g 34 0.94 2.61 1.28 0.00 1.73 2.55 3.40 7.29 _____

```
## Summary for: DW_shoot_g
## # A tibble: 31 × 4
##
     Genotype mean std.dev n missing
##
     <fct>
            <dbl> <dbl> <int>
## 1 EPPN8_H 0.345 0.0744
## 2 EPPN14_H 0.342 0.0710
## 3 EPPN13_H 0.333 0.102
                                   0
## 4 EPPN11_H 0.313 0.0807
                                  1
## 5 EPPN6_H 0.312 0.0617
                                   1
## 6 EPPN9 H 0.311 0.100
                                   0
## 7 EPPN1 H 0.297 0.0715
                                   0
## 8 EPPN5_H 0.286 0.105
                                  0
## 9 EPPN10_H 0.271 0.0633
                                   2
## 10 EPPN9_L 0.265 0.0517
                                  1
## 11 EPPN12_H 0.250 0.0471
                                   3
## 12 EPPN1_L 0.247 0.0759
                                   0
## 13 EPPN3_H 0.245 0.0580
                                  2
## 14 EPPN2 H 0.245 0.0474
                                   2
## 15 EPPN15_H 0.240 0.0473
                                  1
## 16 EPPN3_L 0.239 0.0830
                                   1
## 17 EPPN7_H 0.230 0.0459
                                  1
## 18 EPPN6 L 0.220 0.0343
                                   4
## 19 EPPN5_L 0.209 0.0382
                                   1
## 20 EPPN2 L 0.208 0.0437
                                   3
## 21 EPPN7 L 0.207 0.0279
                                  2
## 22 EPPN8 L 0.207 0.0783
                                   0
## 23 EPPN4 H 0.204 0.0698
                                   2
## 24 EPPN4_L 0.186 0.0313
                                   4
## 25 EPPN20 T 0.177 0.0509
                                   2
## 26 EPPN15_L 0.171 0.0369
## 27 EPPN13_L 0.164 0.0551
                                   1
## 28 EPPN12 L 0.161 0.0583
## 29 EPPN11 L 0.157 0.102
                                   0
## 30 EPPN10 L 0.151 0.0812
## 31 EPPN14_L 0.110 0.0298
                                   1
##
## Summary for: FW shoot g
## # A tibble: 31 × 4
##
     Genotype mean std.dev n_missing
##
     <fct> <dbl>
                     <dbl>
                               <int>
## 1 EPPN14 H 5.42
                                   2
                     0.823
## 2 EPPN8_H 5.23 1.25
                                   3
## 3 EPPN9 H
               4.96
                    1.75
                                   1
## 4 EPPN13_H 4.94
                    1.52
                                   0
               4.94
## 5 EPPN6_H
                    1.13
                                   1
## 6 EPPN11_H 4.62
                    1.45
                                   0
## 7 EPPN1 H 4.56
                    1.06
                                   0
## 8 EPPN5 H 4.32
                    1.63
                                   0
## 9 EPPN3_H
               4.20
                     0.641
                                   4
## 10 EPPN10 H 4.17
                     1.16
                                  1
## 11 EPPN9 L
                                   2
               4.05
                     0.859
## 12 EPPN15 H 3.73
                     0.657
                                   1
## 13 EPPN12 H 3.70
                     0.974
                                   1
## 14 EPPN7 H
               3.56
                     0.744
                                   1
```

```
## 15 EPPN2_H 3.43 0.706
                                  2
## 16 EPPN1 L 3.38 1.32
                                  0
## 17 EPPN3_L 3.32
                   1.59
## 18 EPPN7 L 3.11 0.528
                                  1
## 19 EPPN6_L 3.07
                   0.827
                                  2
## 20 EPPN4 H 2.86 1.21
                                  1
## 21 EPPN2 L 2.73
                   0.356
                                  4
## 22 EPPN20 T 2.56 0.664
                                  1
## 23 EPPN8 L 2.53
                   1.31
## 24 EPPN13 L 2.52 1.11
## 25 EPPN4 L 2.49 0.666
                                  2
## 26 EPPN10_L 2.30
                   1.38
## 27 EPPN15_L 2.20
                                  3
                   0.488
## 28 EPPN12 L 2.09 0.894
                                  0
## 29 EPPN11 L 1.92
                     1.47
                                  0
## 30 EPPN5_L 1.77
                     1.38
                                  0
## 31 EPPN14_L 1.42
                     0.399
                                  1
##
## Summary for: DW_root_g
## # A tibble: 31 × 4
##
     Genotype mean std.dev n_missing
             <dbl> <dbl> <int>
##
     <fct>
## 1 EPPN13_H 0.169 0.0517
## 2 EPPN6_H 0.156 0.0342
## 3 EPPN9 H 0.140 0.0483
## 4 EPPN14_H 0.132 0.0305
## 5 EPPN11_H 0.126 0.0404
## 6 EPPN5_H 0.125 0.0381
## 7 EPPN15 H 0.122 0.0295
## 8 EPPN8_H 0.119 0.0622
## 9 EPPN1 H 0.118 0.0364
## 10 EPPN7_H 0.115 0.0295
## 11 EPPN2 H 0.109 0.0429
## 12 EPPN6 L 0.104 0.0240
                                   3
## 13 EPPN3 H 0.103 0.0363
## 14 EPPN1 L 0.102 0.0219
## 15 EPPN12 H 0.0988 0.0260
                                  3
## 16 EPPN9 L 0.0963 0.0284
## 17 EPPN4_H 0.0947 0.0454
## 18 EPPN7 L 0.0939 0.0131
                                   2
## 19 EPPN3 L 0.0920 0.0455
## 20 EPPN10_H 0.0898 0.0270
                                   1
## 21 EPPN2 L 0.0886 0.0303
                                  1
## 22 EPPN5 L 0.0825 0.0225
## 23 EPPN8 L 0.0770 0.0323
## 24 EPPN13_L 0.0764 0.0349
## 25 EPPN20 T 0.0761 0.0250
                                   1
## 26 EPPN4_L 0.0724 0.0194
                                  2
## 27 EPPN15_L 0.0721 0.0213
                                  1
## 28 EPPN11 L 0.0686 0.0450
## 29 EPPN12 L 0.0588 0.0226
                                   1
## 30 EPPN10_L 0.0477 0.0304
## 31 EPPN14_L 0.0188 0.00501
## Summary for: FW root g
## # A tibble: 31 × 4
```

file:///C:/Users/elise/Documents/Mémoire/Main/Source/Data Analysis/UCL_Data-Analysis.html

```
Genotype mean std.dev n_missing
   <fct> <dbl> <dbl> <int>
## 1 EPPN13_H 4.52 0.984
## 2 EPPN9 H 4.06 1.48
## 3 EPPN6_H 3.96 0.880
## 4 EPPN5 H 3.84 1.51
## 5 EPPN14 H 3.69 0.516
## 6 EPPN11 H 3.63 0.838
## 7 EPPN15_H 3.39 0.757
## 8 EPPN8 H 3.23 1.58
## 9 EPPN3 H 3.12 1.06
## 10 EPPN2_H 3.05 0.595
## 11 EPPN1_H 3.00 0.365
                                7
## 12 EPPN12 H 2.74 0.854
                                1
## 13 EPPN4 H 2.68 1.36
## 14 EPPN7_H 2.52 1.27
                                0
## 15 EPPN9_L 2.50 0.745
                                0
## 16 EPPN10 H 2.34 0.646
                                2
## 17 EPPN3_L 2.27 1.20
                                а
## 18 EPPN7_L 2.17
                   0.156
                                3
## 19 EPPN2_L 2.05 0.550
                                2
## 20 EPPN6_L 2.04 0.756
                                1
## 21 EPPN1_L 2.03 0.776
                                0
## 22 EPPN5_L 2.02 0.810
                                0
## 23 EPPN20 T 1.81 0.570
                                2
## 24 EPPN15_L 1.76 0.682
                                0
## 25 EPPN4_L 1.62 0.380
                                2
## 26 EPPN13_L 1.58 0.796
## 27 EPPN8 L 1.43 0.705
## 28 EPPN11_L 1.20 0.929
## 29 EPPN12 L 1.13 0.545
## 30 EPPN10_L 0.865
                   0.594
## 31 EPPN14 L 0.468
                    0.207
```

2. Exploration of the timeseries data

In this part, we look at the timeseries, S_timeseries and T_timeseries datasets, also using several functions, located in the functions.R script.

Number of data observations per day for the traits of the timeseries datasets

```
print(paste0("No data for", platform))
## [1] "No data forUCL"
```

A. Exploration of the timeseries dataframe

Scatter plots by Genotype

print(paste0("No data for", platform))

[1] "No data forUCL"

Scatterplots for all genotypes by Plant type (Hybride, Line, EPPN20_T) with smooth line.

print(paste0("No data for", platform))

[1] "No data forUCL"

Scatter plots for all genotypes by water treatment

print(paste0("No data for", platform))

[1] "No data forUCL"

B. Exploration of the S timeseries dataframe

Scatter plots by Genotype

print(paste0("No data for", platform))

[1] "No data forUCL"

Scatterplots for all genotypes by Plant type (Hybride, Line, EPPN20 T) with smooth line.

print(paste0("No data for", platform))

[1] "No data forUCL"

Scatter plots for all genotypes by water treatment

print(paste0("No data for", platform))

[1] "No data forUCL"

C. Exploration of the T_timeseries dataframe

Scatter plots by Genotype

print(paste0("No data for", platform))

[1] "No data forUCL"

Scatterplots for all genotypes by Plant type (Hybride, Line, EPPN20_T) with smooth line.

print(paste0("No data for", platform))

[1] "No data forUCL"

Scatter plots for all genotypes by water treatment

print(paste0("No data for", platform))

[1] "No data forUCL"