### WK3

#### 2025-07-02

```
library(tidyverse)
## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr 1.1.4
                      v readr
                                   2.1.5
## v forcats 1.0.0
                       v stringr
                                   1.5.1
## v ggplot2 3.5.2
                       v tibble
                                   3.2.1
## v lubridate 1.9.4
                        v tidyr
                                   1.3.1
## v purrr
              1.0.4
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                 masks stats::lag()
## i Use the conflicted package (<a href="http://conflicted.r-lib.org/">http://conflicted.r-lib.org/</a>) to force all conflicts to become error
library(readxl)
library(lubridate)
library(janitor)
## Attaching package: 'janitor'
## The following objects are masked from 'package:stats':
##
      chisq.test, fisher.test
##
library(purrr)
library(readr)
library(ggthemes)
#setwd("C:/Users/Tobyz/Desktop/S.tort-light-growth/Data")
import\ plant\ data
plant <- read.csv("Data/WL2-2023_Size_Combined.csv") %>%
  clean_names() %>%
  mutate(survey_date = as.Date(survey_date, format = "%m/%d/%Y"))
summary(plant)
   survey_date
                           block
                                            genotype
                                                               pop_mf
## Min.
         :2023-07-03 Length:17336 Length:17336
                                                            Length: 17336
## 1st Qu.:2023-08-02 Class :character Class :character Class :character
## Median:2023-08-30 Mode:character Mode:character Mode:character
## Mean :2023-08-28
## 3rd Qu.:2023-09-20
```

```
##
##
    parent_pop
                                           rep
                                                        height cm
                     Min. : 1.000
                                      Min. : 1.000 Min. : 0.100
## Length:17336
## Class :character
                     1st Qu.: 2.000
                                      1st Qu.: 4.000
                                                      1st Qu.: 1.700
## Mode :character Median : 5.000
                                     Median: 8.000 Median: 3.100
##
                     Mean : 4.584
                                     Mean : 7.932 Mean : 4.491
##
                      3rd Qu.: 6.000
                                      3rd Qu.:11.000
                                                       3rd Qu.: 5.700
##
                      Max.
                           :14.000
                                      Max. :27.000 Max. :39.400
##
                                                       NA's
                                                             :8762
##
    long_leaf_cm
                   survey_notes
## Min.
         :0.100
                   Length: 17336
## 1st Qu.:1.600
                  Class : character
## Median :2.500
                 Mode : character
## Mean
         :2.599
## 3rd Qu.:3.500
## Max.
          :9.000
## NA's
          :9350
consolidate light measurement to a weekly measurement
#import light data
light_raw <- read_csv("Data/IntBioHalfHourTable_clean.txt")</pre>
## Rows: 4063 Columns: 139
## -- Column specification -----
## Delimiter: ","
## dbl (138): RECORD, BattV Max, PTemp C Max, SlrW Avg, SlrW Max, SlrW Min, Sl...
## dttm (1): TIMESTAMP
## i Use 'spec()' to retrieve the full column specification for this data.
## i Specify the column types or set 'show_col_types = FALSE' to quiet this message.
#weekly measurement
weekly_light <- light_raw %>%
 mutate(
   timestamp = ymd_hms(TIMESTAMP),
   SlrW_Avg = as.numeric(SlrW_Avg), # turn into number format
   week = floor_date(timestamp, "week")
 ) %>%
 group_by(week) %>%
 summarise(
   weekly_avg_SlrW = mean(SlrW_Avg, na.rm = TRUE),
    .groups = "drop"
 )
## Warning: There was 1 warning in 'mutate()'.
## i In argument: 'timestamp = ymd_hms(TIMESTAMP)'.
## Caused by warning:
```

Max.

##

:2023-10-20

## ! 84 failed to parse.

```
# result
print(weekly_light)
```

```
## # A tibble: 14 x 2
##
      week
                          weekly_avg_SlrW
##
      <dttm>
## 1 2023-07-30 00:00:00
                                  280.
## 2 2023-08-06 00:00:00
                                  277.
## 3 2023-08-13 00:00:00
                                  186.
## 4 2023-08-20 00:00:00
                                  184.
## 5 2023-08-27 00:00:00
                                  200.
## 6 2023-09-03 00:00:00
                                  211.
## 7 2023-09-10 00:00:00
                                  204.
## 8 2023-09-17 00:00:00
                                  189.
## 9 2023-09-24 00:00:00
                                  159.
## 10 2023-10-01 00:00:00
                                  138.
## 11 2023-10-08 00:00:00
                                  133.
## 12 2023-10-15 00:00:00
                                  135.
## 13 2023-10-22 00:00:00
                                  116.
## 14 NA
                                   -0.616
```

 $compute\ growth\ from\ size\ measurements$ 

```
plant_growth <- plant %>%
  unite("PID", genotype:rep, sep = "_") %>%
  mutate(survey_date = as.Date(survey_date)) %>%
  select(PID, survey_date, height_cm)

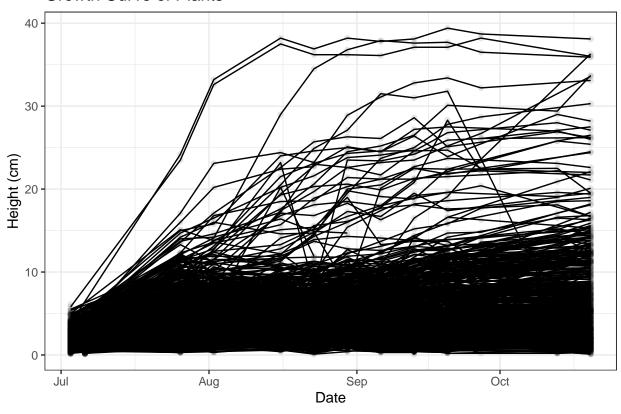
# Will write this data once in a csv and read it in next time
#write_csv(plant_growth, "Data/plant_growth.csv")
#plant_growth = read_csv("Data/plant_growth.csv")

#plot growth curve
ggplot(plant_growth, aes(x = survey_date, y = height_cm, group = PID)) +
  geom_line() +
  geom_point(alpha = 0.1) +
  labs(title = "Growth Curve of Plants", x = "Date", y = "Height (cm)") +
  theme_bw() +
  theme(legend.position = "bottom")
```

## Warning: Removed 8715 rows containing missing values or values outside the scale range
## ('geom\_line()').

## Warning: Removed 8762 rows containing missing values or values outside the scale range ## ('geom\_point()').

#### **Growth Curve of Plants**



```
#check data on these two dates
plant_growth %>%
  filter(survey_date %in% as.Date(c("2023-09-27", "2023-10-13"))) %>%
  group_by(PID, survey_date) %>%
  summarise(has_data = any(!is.na(height_cm)), .groups = "drop") %>%
  pivot_wider(names_from = survey_date, values_from = has_data)
```

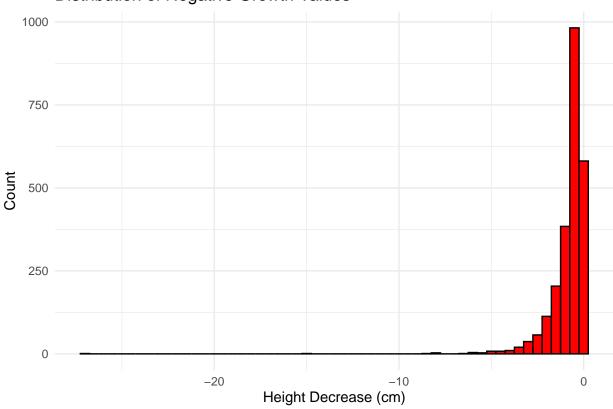
```
## # A tibble: 1,573 x 3
                           '2023-10-13' '2023-09-27'
##
     PID
##
      <chr>
                           <lgl>
                                        <1g1>
## 1 BH_1_10_BH_1_BH_1_10 TRUE
                                        NA
## 2 BH_1_11_BH_1_BH_1_11 FALSE
                                        NA
## 3 BH_1_12_BH_1_BH_1_12 TRUE
                                        NA
## 4 BH_1_13_BH_1_BH_1_13 TRUE
                                        NA
## 5 BH_1_1_BH_1_BH_1_1
                                        NA
                           TRUE
## 6 BH_1_2_BH_1_BH_1_2
                           FALSE
                                        NA
## 7 BH_1_3_BH_1_BH_1_3
                                        FALSE
## 8 BH_1_4_BH_1_BH_1_4
                                        TRUE
                           NA
## 9 BH_1_5_BH_1_BH_1_5
                           NA
                                        FALSE
## 10 BH_1_6_BH_1_BH_1_6
                           NA
                                        FALSE
## # i 1,563 more rows
```

```
plant_growth %>%
  filter(survey_date %in% as.Date(c("2023-09-27", "2023-10-13"))) %>%
  group_by(PID, survey_date) %>%
```

```
summarise(has_data = any(!is.na(height_cm)), .groups = "drop") %>%
  pivot_wider(names_from = survey_date, values_from = has_data) %>%
  mutate(
   status = case when(
      `2023-09-27` == TRUE & `2023-10-13` == TRUE ~ "both",
      `2023-09-27` == TRUE & (`2023-10-13` == FALSE | is.na(`2023-10-13`)) ~ "only 9/27",
      (`2023-09-27` == FALSE | is.na(`2023-09-27`)) & `2023-10-13` == TRUE ~ "only 10/13",
      TRUE ~ "neither"
   )
  ) %>%
  count(status)
## # A tibble: 3 x 2
     status
##
     <chr>
                <int>
## 1 neither
                 1011
## 2 only 10/13
                  336
                  226
## 3 only 9/27
Investigate or filter out plants that show negative growth
#find out plants with negative growth
plant growth %>%
  arrange(PID, survey_date) %>% # arrange in time sequence
  group_by(PID) %>%
 mutate(growth = height_cm - lag(height_cm)) %>% # find out the diff btw nearby dates
  summarise(has negative growth = any(growth < 0, na.rm = TRUE)) %>%
  filter(has_negative_growth) -> neg_growth_plants
neg_growth_plants
## # A tibble: 826 x 2
##
     PID
                           has_negative_growth
##
      <chr>
                           <1g1>
## 1 BH_1_10_BH_1_BH_1_10 TRUE
## 2 BH_1_12_BH_1_BH_1_12 TRUE
## 3 BH 1 13 BH 1 BH 1 13 TRUE
## 4 BH_1_1_BH_1_BH_1_1
                           TRUE
## 5 BH_1_4_BH_1_BH_1_4
                           TRUE
## 6 BH_1_7_BH_1_BH_1_7
                           TRUE
## 7 BH 2 10 BH 2 BH 2 10 TRUE
## 8 BH_2_11_BH_2_BH_2_11 TRUE
## 9 BH 2 12 BH 2 BH 2 12 TRUE
## 10 BH_2_13_BH_2_BH_2_13 TRUE
## # i 816 more rows
#find out tolerance value
neg_growth_values <- plant_growth %>%
 arrange(PID, survey_date) %>%
  group_by(PID) %>%
 mutate(growth = height_cm - lag(height_cm)) %>%
  ungroup() %>%
 filter(growth < 0)</pre>
```

```
ggplot(neg_growth_values, aes(x = growth)) +
  geom_histogram(binwidth = 0.5, fill = "red", color = "black") +
  labs(
    title = "Distribution of Negative Growth Values",
    x = "Height Decrease (cm)",
    y = "Count"
  ) +
  theme_minimal()
```

## Distribution of Negative Growth Values



```
#filter out plants with negative growth < -5
plant_growth_cleaned <- plant_growth_cleaned %>%
    arrange(PID, survey_date) %>%
    group_by(PID) %>%
    mutate(growth = height_cm - lag(height_cm)) %>%
    filter(is.na(growth) | growth >= -5) %>%
    select(-growth) %>%
    ungroup()

check <- plant_growth_cleaned %>%
    arrange(PID, survey_date) %>%
    group_by(PID) %>%
    mutate(growth = height_cm - lag(height_cm)) %>%
    mutate(growth = height_cm - lag(height_cm)) %>%
```

```
filter(growth < -5)

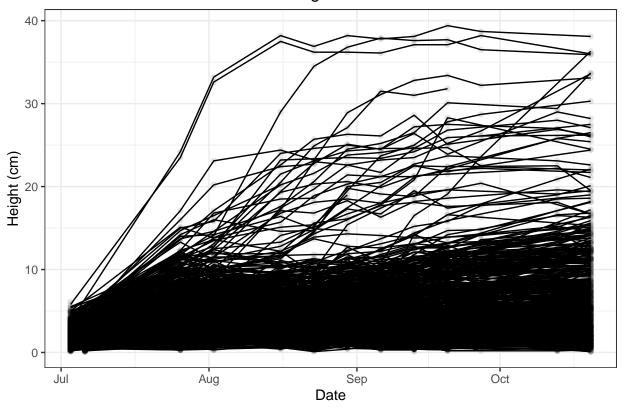
if (nrow(check) == 0) break
}

#plot growth curve
ggplot(plant_growth_cleaned, aes(x = survey_date, y = height_cm, group = PID)) +
    geom_line() +
    geom_point(alpha = 0.1) +
    labs(title = "Growth Curve of Plants without Negative Growth < -5", x = "Date", y = "Height (cm)") +
    theme_bw() +
    theme(legend.position = "bottom")</pre>
```

## Warning: Removed 8715 rows containing missing values or values outside the scale range ## ('geom\_line()').

## Warning: Removed 8762 rows containing missing values or values outside the scale range
## ('geom\_point()').

## Growth Curve of Plants without Negative Growth < -5



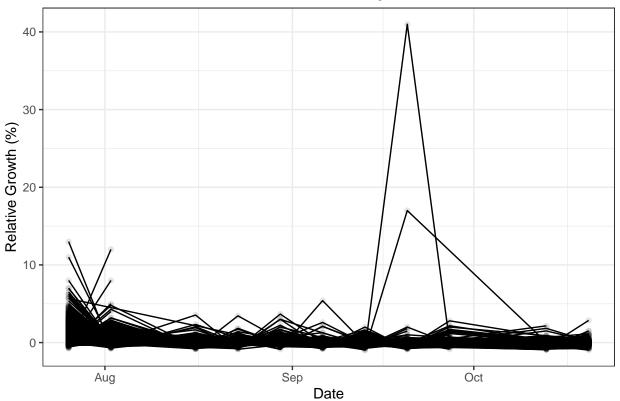
measure growth via relative growth

```
plant_growth_rel <- plant_growth_cleaned %>%
    arrange(PID, survey_date) %>%
    group_by(PID) %>%
    mutate(
```

```
prev_height = lag(height_cm),
    growth = height_cm - prev_height,
   relative_growth = growth / prev_height
  ) %>%
  ungroup()
plant_growth_rel
## # A tibble: 17,312 x 6
     PID
                          survey_date height_cm prev_height growth relative_growth
##
      <chr>
                                         <dbl>
                                                     <dbl> <dbl>
                                                                           <dbl>
                          <date>
## 1 BH_1_10_BH_1_BH_1_10 2023-07-03
                                           2.5
                                                      NA NA
                                           3.1
                                                       2.5 0.6
                                                                          0.24
## 2 BH_1_10_BH_1_BH_1_10 2023-07-26
## 3 BH_1_10_BH_1_BH_1_10 2023-08-02
                                           3.9
                                                       3.1 0.8
                                                                          0.258
## 4 BH_1_10_BH_1_BH_1_10 2023-08-16
                                           2.7
                                                       3.9 -1.2
                                                                         -0.308
## 5 BH_1_10_BH_1_BH_1_10 2023-08-23
                                           3.1
                                                       2.7 0.4
                                                                          0.148
## 6 BH 1 10 BH 1 BH 1 10 2023-08-30
                                           3.1
                                                       3.1 0
                                                                          0
## 7 BH_1_10_BH_1_BH_1_10 2023-09-06
                                           4
                                                       3.1 0.9
                                                                         0.290
## 8 BH_1_10_BH_1_BH_1_10 2023-09-13
                                           5.9
                                                       4 1.9
                                                                          0.475
## 9 BH_1_10_BH_1_BH_1_10 2023-09-20
                                           4.6
                                                      5.9 -1.3
                                                                         -0.220
                                           4.3
                                                       4.6 -0.300
                                                                         -0.0652
## 10 BH_1_10_BH_1_BH_1_10 2023-10-13
## # i 17,302 more rows
plant_growth_rel%>%
  filter(!is.na(relative_growth)) %>%
  ggplot(aes(survey_date, relative_growth, group= PID))+
  geom_line()+
  geom_point(alpha=0.1)+
  labs(title="Relative Growth Curve of Plants without Negative Growth < -5", x = "Date", y = "Relative"
```

theme\_bw()

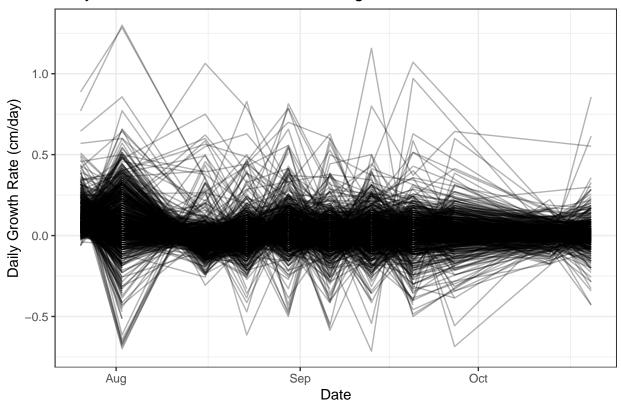




Measure Growth via Daily Growth Rate

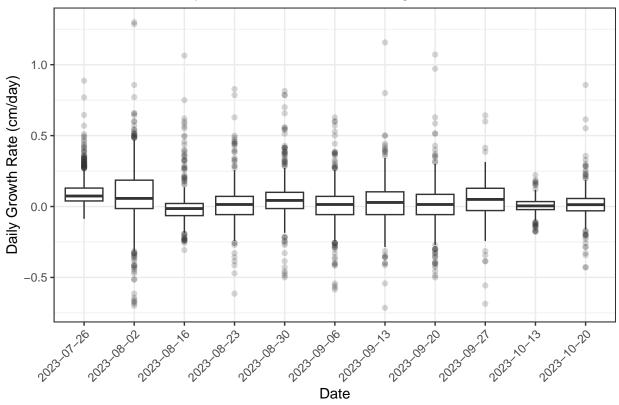
```
#define daily growth rate
plant_growth_daily <- plant_growth_cleaned %>%
  arrange(PID, survey_date) %>%
  group_by(PID) %>%
  mutate(
   prev_height = lag(height_cm),
   prev_date = lag(survey_date),
    days_elapsed = as.numeric(survey_date - prev_date),
    daily_growth = (height_cm - prev_height) / days_elapsed
  ) %>%
  ungroup()
#Line graph
plant_growth_daily %>%
  filter(!is.na(daily_growth)) %>%
  ggplot(aes(x = survey_date, y = daily_growth, group = PID)) +
  geom_line(alpha = 0.3) +
  labs(title = "Daily Growth Rate of Plants without Negative Growth < -5", y = "Daily Growth Rate (cm/d
  theme_bw()
```





```
#boxplot
plant_growth_daily %>%
  filter(!is.na(daily_growth), days_elapsed > 0) %>%
  mutate(date_factor = factor(survey_date, levels = sort(unique(survey_date))))%>%
  ggplot(aes(x = date_factor, y = daily_growth)) +
  geom_boxplot(outlier.alpha = 0.2) +
  labs(title = "Distribution of Daily Growth Rates without Negative Growth < -5", x = "Date", y = "Dail'
  theme_bw()+
  theme(axis.text.x = element_text(angle = 45, hjust = 1))</pre>
```





Correlate Growth with Solar Radiation

```
#Align plant growth data to week
plant_weekly <- plant_growth_daily %>%
  filter(!is.na(daily_growth), days_elapsed > 0) %>%
  mutate(week = floor_date(survey_date, "week"))

#Adds `weekly_avg_SlrW` to plant data
plant_with_light <- plant_weekly %>%
  left_join(weekly_light, by = "week")

#Calculate correlation
cor_result <- cor(
  plant_with_light$daily_growth,
  plant_with_light$weekly_avg_SlrW,
use = "complete.obs"
)

print(cor_result)</pre>
```

```
## [1] 0.1299395
```

```
ggplot(plant_with_light, aes(x = weekly_avg_SlrW, y = daily_growth)) +
geom_point(alpha = 0.2) +
geom_smooth(method = "lm", color = "red") +
```

```
labs(title = "Correlation between Light and Growth", x = "Weekly Avg Light (SlrW)", y = "Daily Growth
theme_bw()
```

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

## Warning: Removed 1069 rows containing non-finite outside the scale range
## ('stat_smooth()').

## Warning: Removed 1069 rows containing missing values or values outside the scale range
## ('geom_point()').
```

# Correlation between Light and Growth

