

Proj

Mānuka background(cultural,uses,industry)

Experimental Cross Design

Two east cape parents, F1 cross siblings

F1 Layout Map indicates the layout of the F1 generation resulting from a cross between EC201 and EC103 parents. Column 1 is approximately lengthways facing north.

Height Diagram Diagram indicates the areas of leaf collection regarding height. Each Tree generally had 10 leaves collected (some trees have been sampled more than once), leaves were selected from low, mid and high points of the tree. The first leaf sampled from each tree was measured twice for replication comparison.

Import and Arrange Data Dualex was used to measure transmittance through all samples included in this dataset, more information on how this works can be found in Clara Rey Caramés, 2015.

"Full.xlsx"

- Surface content of chlorophyll in g/cm^2 (Chl)
- Epidermal Flavoid content in absorbance units(Au); Flavonol(Flav) and Anthocyanin(Anth)
- Nitrogen Balance Index status is calculated using Chlorophyll and Flavonol values automatically (NBI)

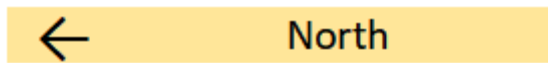
It also contains information about the block position, the leaf height information, and presense or absence of flowering

Sheet "Dup" contains only the replicated samples

Checking Data Quality from Replicate Leaf Measures. Replicates exist in two groups; first leaf measurement (Y1), and the second leaf measurement (Y2). The second measurement was taken after removing the leaf following the first reading, to help establish the presence of any leaf positional bias.

Replicate Plots and Data Overview

##	Rep.	AnthMean	AnthSD	ChlMean	ChlSD	FlavMean	FlavSD
## 1	Y1	0.09830189	0.04539806	34.93672	14.55641	1.979969	0.2863768
## 2	Y2	0.09768153	0.04439018	36.59920	14.51963	1.954280	0.2796720



Row	Column 1	Column 2	Column 3	Column 4
50	IN4DF	IN4F4	IN4GN	IN4GP
49	IN4DE	IN4F3	IN4GM	IN4G5
48	IN4DD	IN4F2	IN4GL	
47	IN4DC	IN4F1	IN4GK	IN4GV
46	IN4DA	IN4F0	IN4GJ	IN4GW
45	IN4D9	IN4E2	IN4GH	IN4GX
44	IN4D8	IN4EY	IN4GG	
43	IN4D7	IN4EX	IN4GF	
42	IN4D6	IN4EW	IN4GE	
41	IN4D5	IN4EV	IN4GD	IN4H1
40	IN4D4	IN4ET	IN4GC	
39	IN4D3	IN4ES	IN4GB	
38	IN4D2	IN4ER	IN4GA	
37	IN4D0	IN4EP	IN4G9	
36	IN4CY	IN4EN	IN4G8	
35	IN4CX	IN4EM	IN4G7	IN4H7
34	IN4CW	IN4EL	IN4G6	IN4H8
33	IN4CV	IN4EK	IN4G5	
32	IN4CT	IN4EJ	IN4G4	IN4HA
31	IN4CS	IN4EH	IN4G3	
30	IN4CR	IN4EG	IN4G2	
29	IN4CP	IN4EF	IN4G1	
28	IN4CN	IN4EE	IN4G0	
27	IN4CM	IN4ED	IN4F2	
26	IN4CL	IN4EC	IN4FY	
25	IN4CK	IN4EB	IN4FX	
24	IN4CJ	IN4EA	IN4FW	
23	IN4CH	IN4E9	IN4FV	IN4HK
22	IN4CG	IN4E8	IN4FT	
21	IN4CE	IN4E6	IN4FS	
20	IN4CD	IN4E4	IN4FR	
19	IN4CC	IN4E3	IN4FP	
18	IN4CB	IN4E2	IN4FN	
17		IN4E1	IN4FM	
16	IN4C9	IN4E0	IN4FL	
15	IN4C8	IN4DZ	IN4FK	
14	IN4C7	IN4DY	IN4FJ	
13	IN4C6	IN4DX	IN4FH	
12	IN4C5	IN4DW	IN4FG	
11	IN4C4	IN4DV	IN4FF	
10	IN4C3	IN4DT	IN4FE	
9	IN4C2	IN4DFS	IN4FD	
8	IN4C1	IN4DR	IN4FC	
7	IN4C0	IN4DP	IN4FB	
6	IN4B2	IN4DM	IN4FA	
5	IN4BY	IN4DL	IN4F9	
4	IN4BX	IN4DK	IN4F8	
3	IN4BW	IN4DJ	IN4F7	
2	IN4BV	IN4DH	IN4F6	
1	IN4BT	IN4DG	IN4F5	

Figure 1: “Map of F1 Trees”

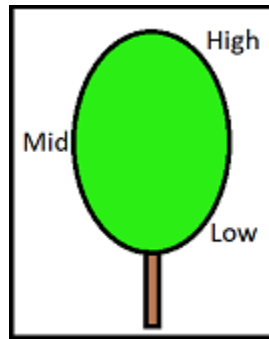


Figure 2: “Diagram of leaf collection levels”

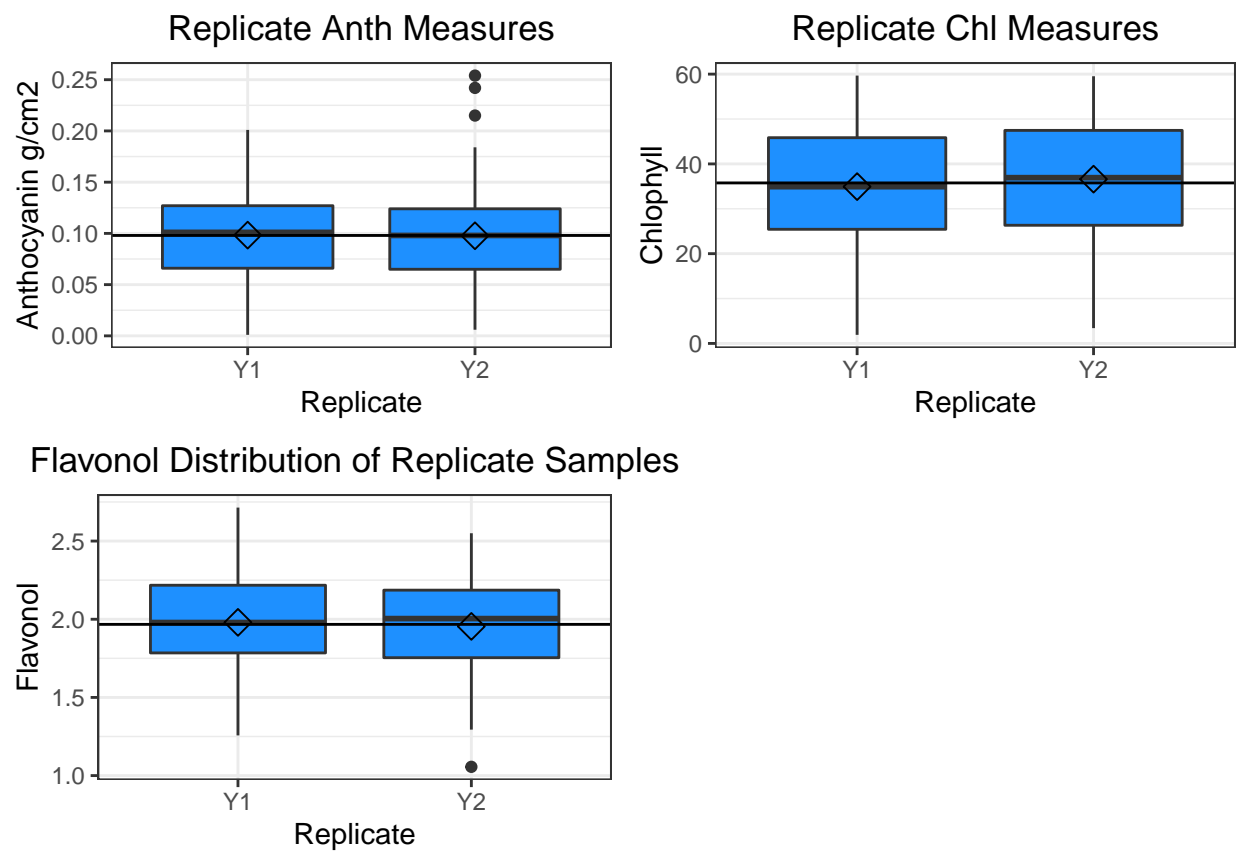


Figure 3: Plots of replicate measure distributions, medians (lines), means (empty diamonds)

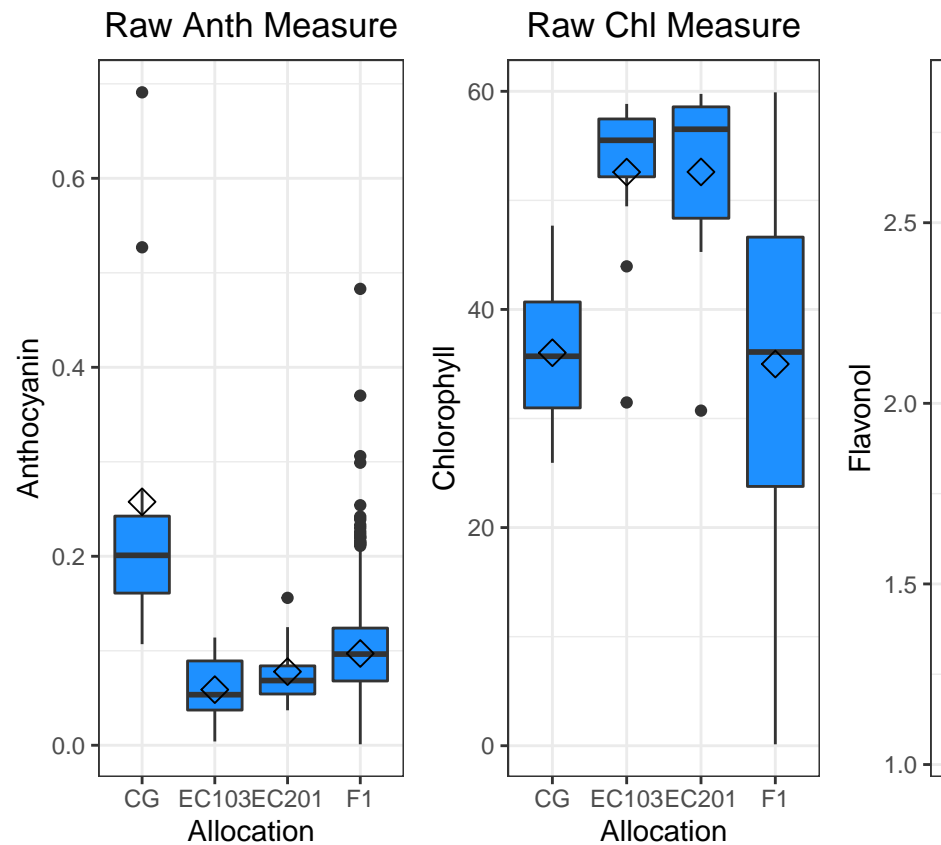
Replicate ANOVAs

```
## Analysis of Variance Table
##
## Response: Dup$Anth
##           Df Sum Sq Mean Sq F value Pr(>F)
## Dup$Rep.    1  0.00003 0.0000304  0.0151 0.9023
## Residuals  314  0.63303 0.0020160
```

```
## Analysis of Variance Table
##
## Response: Dup$Chl
##           Df Sum Sq Mean Sq F value Pr(>F)
## Dup$Rep.    1    218   218.33   1.033 0.3102
## Residuals  314  66366   211.36
```

```
## Analysis of Variance Table
##
## Response: Dup$Flav
##           Df Sum Sq Mean Sq F value Pr(>F)
## Dup$Rep.    1  0.0521  0.052129  0.6506 0.4205
## Residuals  314 25.1596  0.080126
```

Replicate measures and plots appear to correspond well - indicative that the measurements from the dualux are reliable. Might pay to remove the outliers.



Plots of leaf chemicals across groups

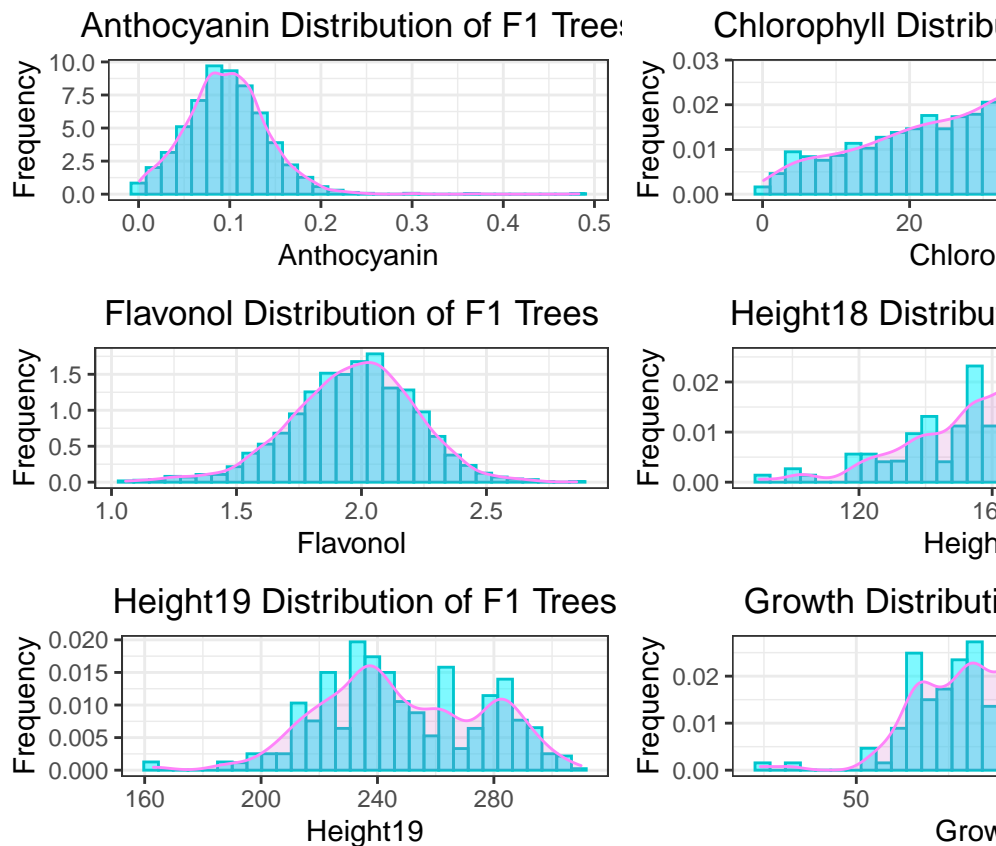
Data Summary

##	Allocation	AnthMean	AnthSD	ChlMean	ChlSD	FlavMean	FlavSD
## 1	CG	0.25763636	0.18306079	36.02836	7.090510	2.169455	0.2021457
## 2	EC103	0.05883333	0.03576650	52.59242	7.889391	1.599833	0.2334250
## 3	EC201	0.07790000	0.03665288	52.60670	9.291990	1.710800	0.2217590
## 4	F1	0.09711830	0.04437066	34.99601	15.144271	1.967003	0.2479930

Anthocyanin Crimson Glory(CG) outgroup apparent, F1 population more similar to parental trees.

Chlorophyll F1 very diverse, overlaps with both parent trees and outgroup, more similar median and distribution to outgroup.

Flavonol F1 overlaps with parents and outgroup, slightly more similar to parental trees.



Plots of unadjusted F1 chemical

Anthocyanin left bias, long tail

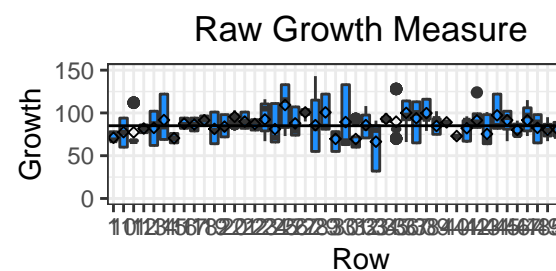
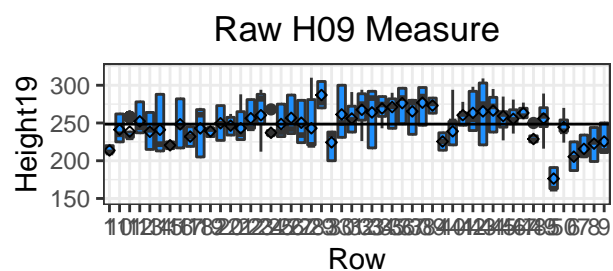
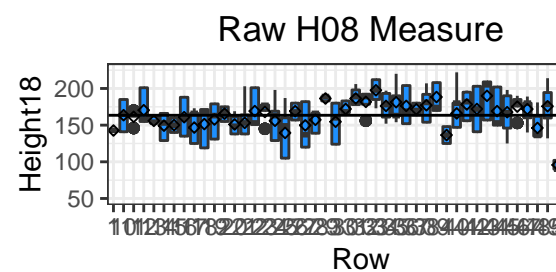
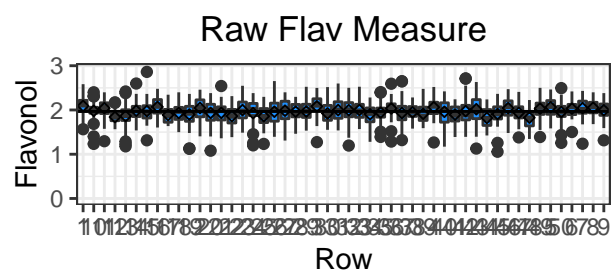
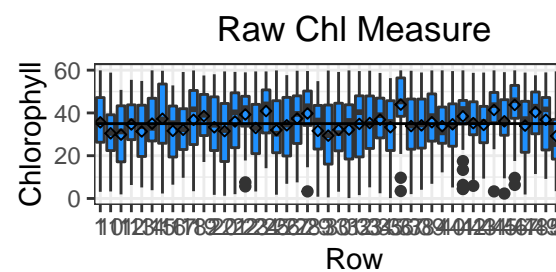
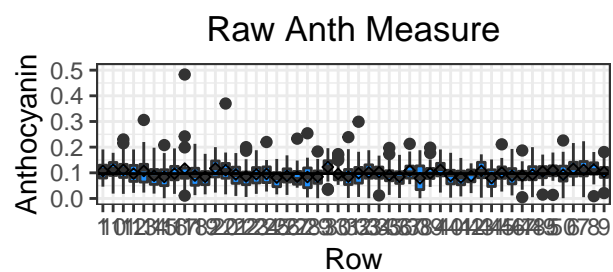
Chlorophyll right bias, truncated

Flavonol typical bell curve with slightly long tails both directions

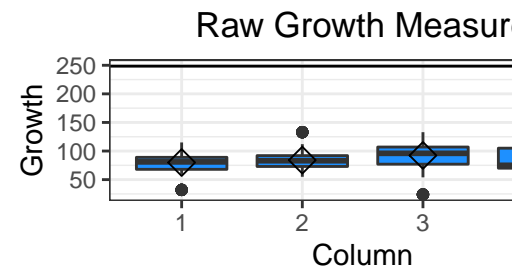
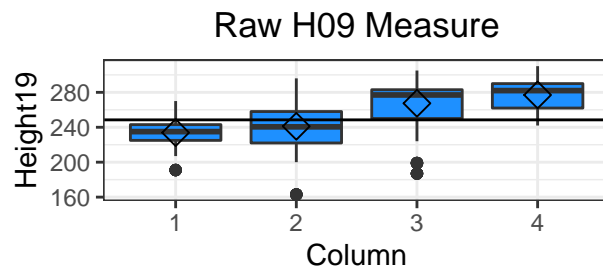
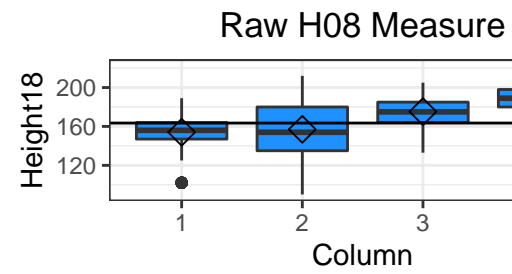
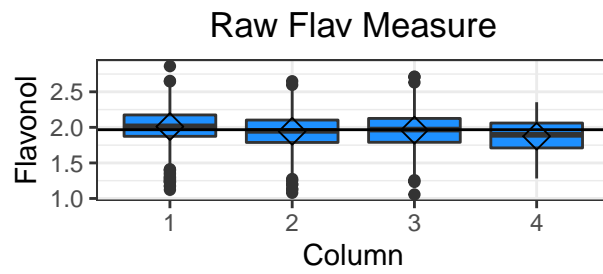
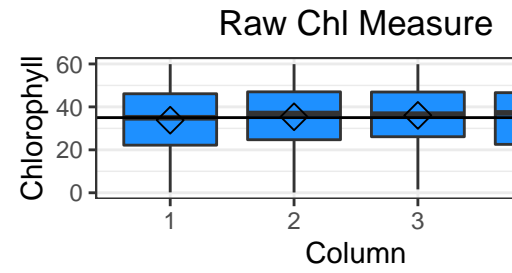
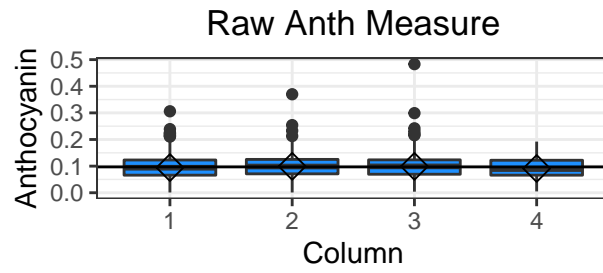
Height 2018 lumpy normal distribution

Height 2019 appears to be two overlapping distributions, maybe 3

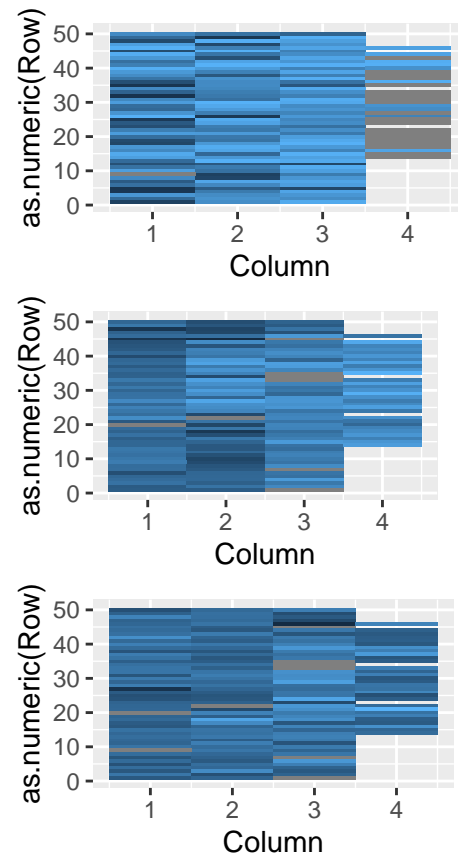
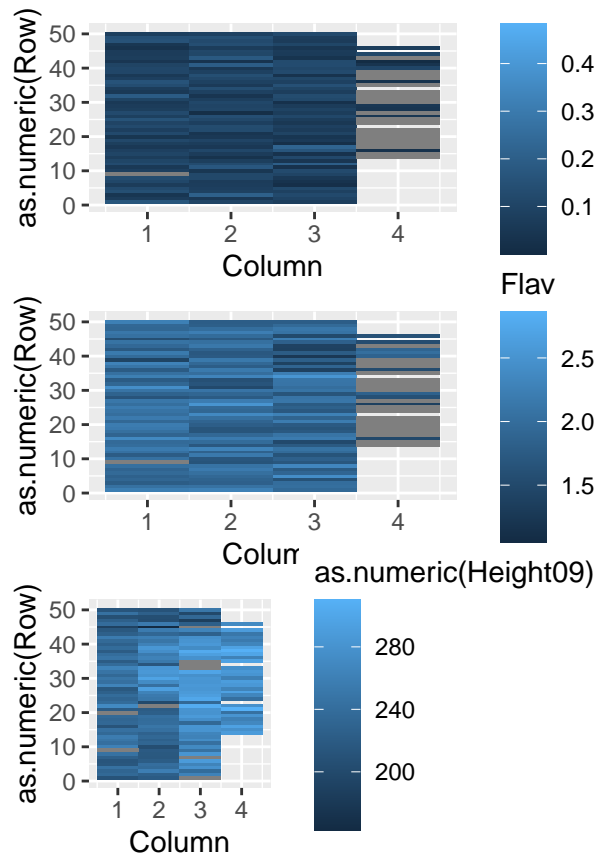
Growth maybe 3 overlapping populations



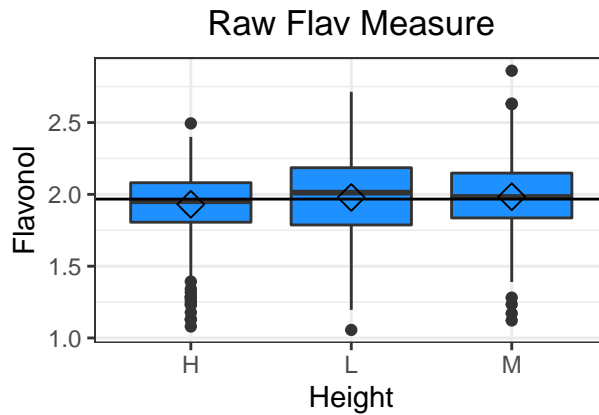
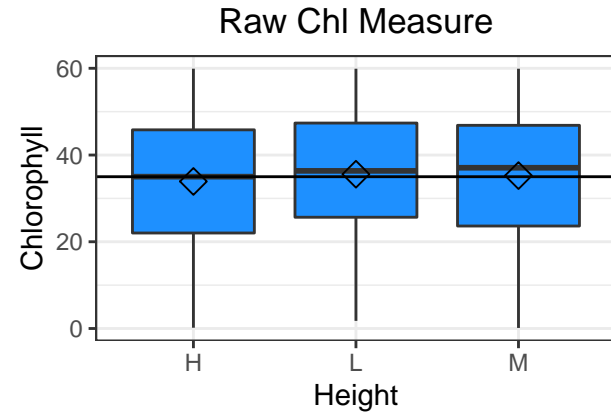
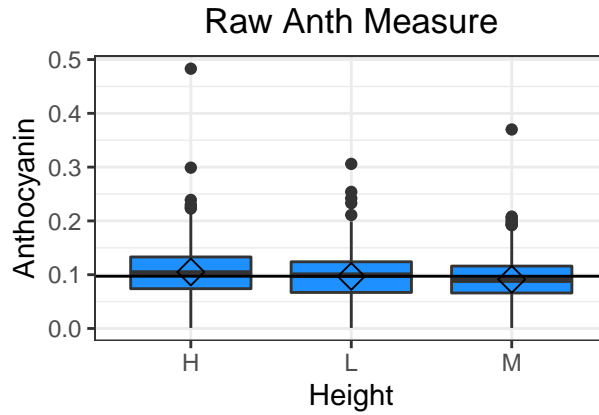
Row Comparisons



Column Comparisons



Row v Column Heatmaps



Height Plots

ANOVAs comparing Height, Row and Columns for each chemical

```
## Analysis of Variance Table
##
## Response: F1$Anth
##          Df Sum Sq Mean Sq F value    Pr(>F)
## F1$Row      49  0.2346  0.0047878  2.5717 2.467e-08 ***
## F1$Column    1  0.0020  0.0019565  1.0509  0.3054
## F1$Height    2  0.0520  0.0260053 13.9687 9.587e-07 ***
## Residuals 1739  3.2375  0.0018617
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
## Analysis of Variance Table
##
## Response: F1$Chl
##          Df Sum Sq Mean Sq F value    Pr(>F)
## F1$Row      49 21394  436.61  1.9560 0.0001012 ***
## F1$Column    1   470  469.95  2.1054 0.1469578
## F1$Height    2   737  368.69  1.6517 0.1920166
## Residuals 1739 388163  223.21
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
## Analysis of Variance Table
```

```
##
## Response: F1$Flav
##           Df Sum Sq Mean Sq F value    Pr(>F)
## F1$Row      49  8.215  0.16765   2.9272 1.204e-10 ***
## F1$Column    1  1.379  1.37854  24.0698 1.016e-06 ***
## F1$Height    2  0.957  0.47848   8.3545 0.0002449 ***
## Residuals 1739 99.597  0.05727
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Analysis of Variance Table

```
##
## Response: F1$Height08
##           Df Sum Sq Mean Sq F value    Pr(>F)
## F1$Row      49 493101   10063  42.056 < 2.2e-16 ***
## F1$Column    1 154439  154439 645.422 < 2.2e-16 ***
## Residuals 1673 400321     239
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Analysis of Variance Table

```
##
## Response: as.numeric(F1$Height09)
##           Df Sum Sq Mean Sq F value    Pr(>F)
## F1$Row      49 626751   12791  51.649 < 2.2e-16 ***
## F1$Column    1 275536  275536 1112.606 < 2.2e-16 ***
## Residuals 1672 414069     248
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Analysis of Variance Table

```
##
## Response: as.numeric(F1$Growth)
##           Df Sum Sq Mean Sq F value    Pr(>F)
## F1$Row      49 145650   2972.5  12.725 < 2.2e-16 ***
## F1$Column    1  17049 17049.2   72.985 < 2.2e-16 ***
## Residuals 1672 390575     233.6
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Anthocyanin Significant Row and Height effects

Chlorophyll Significant Row effect

Flavonol Significant row,column and height effects

Height 2018 Very Significant effect of row and column

Height 2019 Very Significant effect of row and column

Growth difference (Height 2019- Height 2018) Very Significant effect of row and column - need to double check might be from later in analysis

Adjusting Row, Column and Height measures

```
##      Height Column Row Tree.ID Allocation Rep. measure Flower Flower.Level
## 490      H      3   9   IN4FM      F1      N      11      N      0
## 1307     M      2  13   IN4E4      F1      N      7      N      0
##      Chl  Flav  Anth  NBI Height08 Height09 Growth AnthRowMeans ChlRowMeans
## 490 59.122 1.640 0.483 36.05      166      245      79      0.1163182 32.12650
## 1307 18.511 1.401 0.370 13.21      138      234      96      0.1105227 31.51561
##      FlawRowMeans H18RowMeans H19RowMeans AnthColMeans ChlColMeans FlavColMeans
## 490      1.886773      146.6957      232.0 0.09897588 36.04402 1.960649
## 1307      1.933068      151.0000      246.5 0.09924281 35.39257 1.944405
##      H18ColMeans H19ColMeans AnthHeightMeans ChlHeightMeans FlavHeightMeans
## 490      175.1734      267.4482      0.10500200      33.90735      1.930788
## 1307      157.0018      241.0717      0.09133645      35.25540      1.983505
##      AdjAnth
## 490 0.4559164
## 1307 0.3623774
```

Analysis of Variance Table

##

Response: F1\$AdjAnth

```
##      Df Sum Sq Mean Sq F value Pr(>F)
## F1$Row      1 0.00026 0.00025836 0.1518 0.6969
## F1$Column    1 0.00118 0.00118033 0.6934 0.4051
## F1$Height    2 0.00018 0.00008970 0.0527 0.9487
## Residuals 1785 3.03861 0.00170230
```

Analysis of Variance Table

##

Response: F1\$AdjChl

```
##      Df Sum Sq Mean Sq F value Pr(>F)
## F1$Row      1      1      0.89 0.0041 0.9489
## F1$Column    1     72     72.41 0.3349 0.5628
## F1$Height    2    789    394.51 1.8249 0.1615
## Residuals 1785 385885 216.18
```

Analysis of Variance Table

##

Response: F1\$AdjFlav

```
##      Df Sum Sq Mean Sq F value Pr(>F)
## F1$Row      1 0.019 0.018555 0.3357 0.5624
## F1$Column    1 0.007 0.007400 0.1339 0.7145
## F1$Height    2 0.001 0.000454 0.0082 0.9918
## Residuals 1785 98.652 0.055267
```

Analysis of Variance Table

##

Response: F1\$AdjH18

```
##      Df Sum Sq Mean Sq F value Pr(>F)
## F1$Row      1 2625 2625.3 11.005 0.0009277 ***
## F1$Column    1 6285 6285.4 26.347 3.176e-07 ***
## Residuals 1719 410084 238.6
```

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```
## Analysis of Variance Table
##
## Response: F1$AdjH19
##           Df Sum Sq Mean Sq F value    Pr(>F)
## F1$Row      1   3064   3063.9   12.788 0.0003586 ***
## F1$Column    1  14757  14756.5   61.589 7.385e-15 ***
## Residuals 1718 411630    239.6
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
## Analysis of Variance Table
##
## Response: as.numeric(F1$Growth)
##           Df Sum Sq Mean Sq F value    Pr(>F)
## F1$Row      1     26    26.07  0.1178 0.731482
## F1$Column    1  1852 1851.64   8.3672 0.003869 **
## Residuals 1718 380187    221.30
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Adjust height row and column effects

```
## Analysis of Variance Table
##
## Response: F1$AdjH182
##           Df Sum Sq Mean Sq F value    Pr(>F)
## F1$Row      1    174 173.971   0.7742 0.3790
## F1$Column    1     7   6.782   0.0302 0.8621
## Residuals 1718 386029 224.697
```

```
## Analysis of Variance Table
##
## Response: F1$AdjH192
##           Df Sum Sq Mean Sq F value    Pr(>F)
## F1$Row      1    423  423.06   1.8915 0.1692147
## F1$Column    1  3107 3107.07  13.8915 0.0001999 ***
## Residuals 1718 384260    223.67
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
## Analysis of Variance Table
##
## Response: as.numeric(F1$Growth)
##           Df Sum Sq Mean Sq F value    Pr(>F)
## F1$Row      1    54.44   54.44   39.712 3.731e-10 ***
## F1$Column    1 2823.51 2823.51 2059.493 < 2.2e-16 ***
## Residuals 1718 2355.33     1.37
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
## Analysis of Variance Table
##
```

```
## Response: F1$AdjH193
##           Df Sum Sq Mean Sq F value Pr(>F)
## F1$Row      1     63  63.269   0.2839 0.5942
## F1$Column    1    210 209.574   0.9403 0.3323
## Residuals 1718 382892 222.871

## Analysis of Variance Table
##
## Response: as.numeric(F1$Growth)
##           Df Sum Sq Mean Sq F value    Pr(>F)
## F1$Row      1   54.44    54.44   39.712 3.731e-10 ***
## F1$Column    1 2823.51 2823.51 2059.493 < 2.2e-16 ***
## Residuals 1718 2355.33    1.37
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

