

Untitled3

Telha H. Rehman

R Markdown

This is an R Markdown document. Markdown is a simple formatting syntax for authoring HTML, PDF, and MS Word documents. For more details on using R Markdown see <http://rmarkdown.rstudio.com>.

When you click the **Knit** button a document will be generated that includes both content as well as the output of any embedded R code chunks within the document. You can embed an R code chunk like this:

```
Sys.time()

## [1] "2022-03-10 11:30:37 PST"

library(tinytex)
library(tidyverse)
library(cowplot)
library(Cairo)
library(modelr)
library(gridExtra)
library(mixtools)
library(nlme)
library(car)
library(emmeans)
library(MuMIn)
library(ggpmisc)
library(gtable)
library(grid)
library(RColorBrewer)
library(segmented)
library(data.table)
library(scales)
library(sm)
library(rcompanion)
library(nlstools)
```

DATA

GreenSeeker NDVI Data

```
gs_ndvi_data <- read_csv(file = "DATA/PI_greenseeker_data.csv")

## Rows: 328 Columns: 17
## -- Column specification -----
## Delimiter: ","
## chr (5): site_year, NDVI_1, NDVI_2, NDVI_3, NDVI_4
## dbl (12): year, exp_plot_number, block, plot, N_level, N_level_kgha, biomass...
```

```
##
## 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.
str(gs_ndvi_data , give.attr = FALSE)

## spec_tbl_df [328 x 17] (S3: spec_tbl_df/tbl_df/tbl/data.frame)
## $ site_year      : chr [1:328] "Arbuckle-15" "Arbuckle-15" "Arbuckle-15" "Arbuckle-15" ...
## $ year           : num [1:328] 2015 2015 2015 2015 2015 ...
## $ exp_plot_number : num [1:328] 101 102 103 104 105 201 202 203 204 205 ...
## $ block          : num [1:328] 1 1 1 1 1 2 2 2 2 2 ...
## $ plot           : num [1:328] 1 2 3 4 5 1 2 3 4 5 ...
## $ N_level        : num [1:328] 125 225 0 75 175 0 175 125 225 75 ...
## $ N_level_kgha    : num [1:328] 125 225 0 75 175 0 175 125 225 75 ...
## $ biomass_plus_bag_g: num [1:328] 414 472 281 386 455 304 402 322 418 336 ...
## $ paper_bag_g     : num [1:328] 45 45 45 45 45 45 45 45 45 45 ...
## $ ring_size       : num [1:328] 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 ...
## $ num_of_paper_bags : num [1:328] 1 1 1 1 1 1 1 1 1 1 ...
## $ sample_weight_mg : num [1:328] 4.84 5.12 4.78 5.15 4.93 ...
## $ sample_N_ug     : num [1:328] 117.1 153.4 64.9 92.9 116 ...
## $ NDVI_1          : chr [1:328] "0.77" "0.82" "0.56" "0.72" ...
## $ NDVI_2          : chr [1:328] "n/a" "n/a" "n/a" "n/a" ...
## $ NDVI_3          : chr [1:328] "n/a" "n/a" "n/a" "n/a" ...
## $ NDVI_4          : chr [1:328] "n/a" "n/a" "n/a" "n/a" ...

gs_ndvi_data <- gs_ndvi_data %>%
  filter(!year %in% c("2015" , "2016") & N_level_kgha != 275) #remove the years we don't need for this

str(gs_ndvi_data , give.attr = FALSE)

## spec_tbl_df [240 x 17] (S3: spec_tbl_df/tbl_df/tbl/data.frame)
## $ site_year      : chr [1:240] "Nicolaus-17" "Nicolaus-17" "Nicolaus-17" "Nicolaus-17" ...
## $ year           : num [1:240] 2017 2017 2017 2017 2017 ...
## $ exp_plot_number : num [1:240] 101 102 103 104 106 107 202 203 204 205 ...
## $ block          : num [1:240] 1 1 1 1 1 1 2 2 2 2 ...
## $ plot           : num [1:240] 1 2 3 4 6 7 2 3 4 5 ...
## $ N_level        : num [1:240] 225 0 125 175 45 75 75 125 0 175 ...
## $ N_level_kgha    : num [1:240] 225 0 125 175 45 75 75 125 0 175 ...
## $ biomass_plus_bag_g: num [1:240] 361 264 318 360 285 ...
## $ paper_bag_g     : num [1:240] 45.5 45.5 45.5 45.5 45.5 45.5 45.5 45.5 45.5 45.5 ...
## $ ring_size       : num [1:240] 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 ...
## $ num_of_paper_bags : num [1:240] 1 1 1 1 1 1 1 1 1 1 ...
## $ sample_weight_mg : num [1:240] 3.17 3.42 3.09 3.07 3.2 ...
## $ sample_N_ug     : num [1:240] 79.4 61.1 80.7 95.9 63.6 ...
## $ NDVI_1          : chr [1:240] "0.77" "0.50" "0.67" "0.79" ...
## $ NDVI_2          : chr [1:240] "0.77" "0.52" "0.67" "0.78" ...
## $ NDVI_3          : chr [1:240] "0.79" "0.46" "0.71" "0.79" ...
## $ NDVI_4          : chr [1:240] "n/a" "n/a" "n/a" "n/a" ...

gs_ndvi_data <- gs_ndvi_data[c(1:240), c(1:17)] #removes the empty rows and columns from the data frame

gs_ndvi_data$block <- factor(gs_ndvi_data$block)
gs_ndvi_data$year <- factor(gs_ndvi_data$year)
gs_ndvi_data$plot <- factor(gs_ndvi_data$plot)
gs_ndvi_data$N_level_kgha_f <- factor(gs_ndvi_data$N_level_kgha)
gs_ndvi_data$exp_plot_number <- factor(gs_ndvi_data$exp_plot_number)
```

```

gs_ndvi_data$site_year <- factor(gs_ndvi_data$site_year , levels = c( "Nicolaus-17" , "Williams-17" , "
gs_ndvi_data$NDVI_1 <- as.numeric(as.character(gs_ndvi_data$NDVI_1))

## Warning: NAs introduced by coercion

gs_ndvi_data$NDVI_2 <- as.numeric(as.character(gs_ndvi_data$NDVI_2))

## Warning: NAs introduced by coercion

gs_ndvi_data$NDVI_3 <- as.numeric(as.character(gs_ndvi_data$NDVI_3))
gs_ndvi_data$NDVI_4 <- as.numeric(as.character(gs_ndvi_data$NDVI_4)) #gets the data right

## Warning: NAs introduced by coercion

str(gs_ndvi_data , give.attr = FALSE)

## tibble [240 x 18] (S3: tbl_df/tbl/data.frame)
## $ site_year      : Factor w/ 10 levels "Nicolaus-17",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ year           : Factor w/ 3 levels "2017","2018",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ exp_plot_number : Factor w/ 27 levels "101","102","103",...: 1 2 3 4 6 7 9 10 11 12 ...
## $ block          : Factor w/ 4 levels "1","2","3","4": 1 1 1 1 1 1 2 2 2 2 ...
## $ plot           : Factor w/ 7 levels "1","2","3","4",...: 1 2 3 4 6 7 2 3 4 5 ...
## $ N_level         : num [1:240] 225 0 125 175 45 75 75 125 0 175 ...
## $ N_level_kgha    : num [1:240] 225 0 125 175 45 75 75 125 0 175 ...
## $ biomass_plus_bag_g: num [1:240] 361 264 318 360 285 ...
## $ paper_bag_g     : num [1:240] 45.5 45.5 45.5 45.5 45.5 45.5 45.5 45.5 45.5 45.5 ...
## $ ring_size       : num [1:240] 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 ...
## $ num_of_paper_bags : num [1:240] 1 1 1 1 1 1 1 1 1 1 ...
## $ sample_weight_mg : num [1:240] 3.17 3.42 3.09 3.07 3.2 ...
## $ sample_N_ug      : num [1:240] 79.4 61.1 80.7 95.9 63.6 ...
## $ NDVI_1           : num [1:240] 0.77 0.5 0.67 0.79 0.68 0.63 0.66 0.68 0.54 0.77 ...
## $ NDVI_2           : num [1:240] 0.77 0.52 0.67 0.78 0.69 0.56 0.65 0.68 0.5 0.75 ...
## $ NDVI_3           : num [1:240] 0.79 0.46 0.71 0.79 0.68 0.53 0.63 0.64 0.5 0.77 ...
## $ NDVI_4           : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ N_level_kgha_f   : Factor w/ 11 levels "0","45","75",...: 10 1 5 8 2 3 3 5 1 8 ...

gs_ndvi_data <- gs_ndvi_data %>%
  mutate( biomass_dry_wt = biomass_plus_bag_g - (paper_bag_g * num_of_paper_bags) ,
          aboveground_biomass = (biomass_dry_wt / ring_size) * 10 ,
          n_content = sample_N_ug / sample_weight_mg ,
          PI_N_Uptake = (aboveground_biomass * n_content) / 1000 #n uptake in kg per ha
          )#processes the data

gs_ndvi_data <- gs_ndvi_data %>%
  rowwise() %>%
  mutate(NDVI = mean(c( NDVI_1 , NDVI_2 , NDVI_3 , NDVI_4 ) , na.rm = T)) #takes average of four NDVI re

gs_ndvi_data <- dplyr::select(gs_ndvi_data ,
                             site_year,
                             year,
                             exp_plot_number,
                             block,
                             plot,
                             N_level_kgha,
                             aboveground_biomass,

```

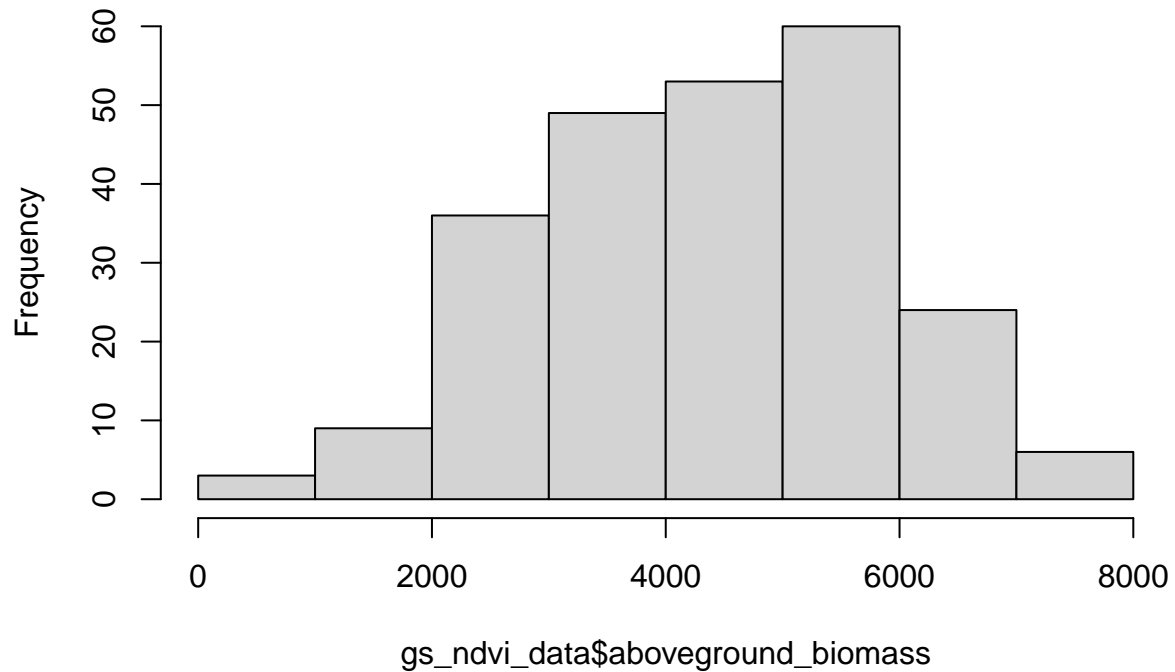
```

n_content,
PI_N_Uptake,
NDVI) #selects the relevant columns

gs_ndvi_data$site_year <- factor(gs_ndvi_data$site_year , levels = c( "Nicolaus-17" , "Williams-17" , "
hist(gs_ndvi_data$aboveground_biomass)

```

Histogram of gs_ndvi_data\$aboveground_biomass

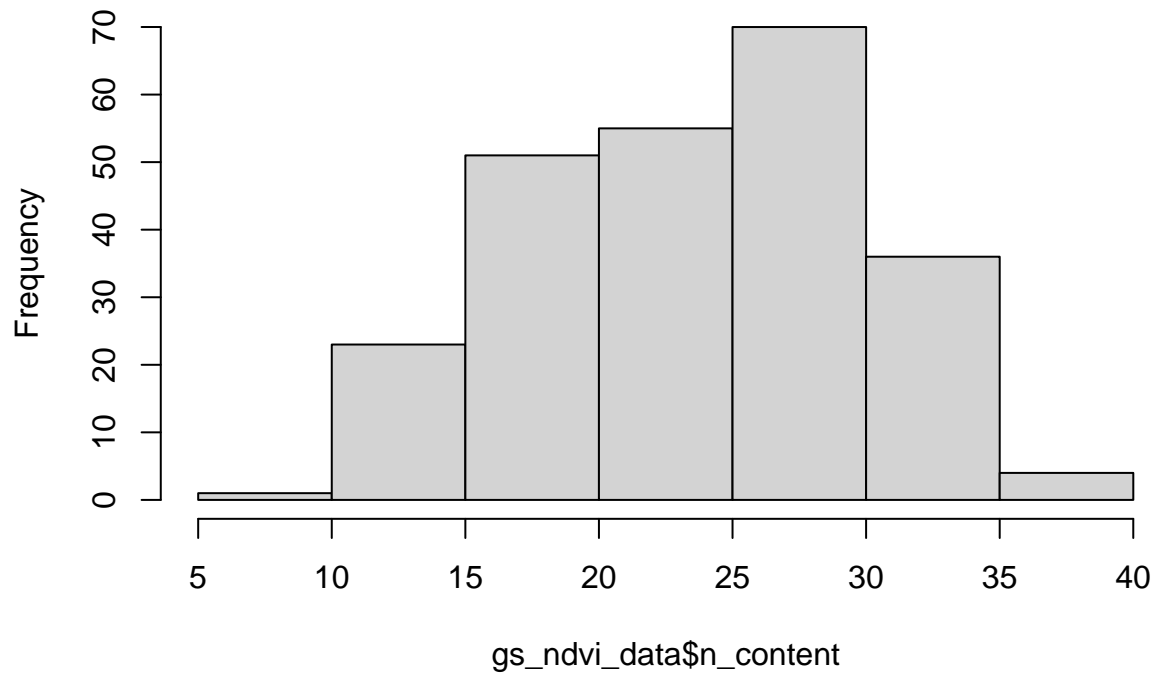


```

hist(gs_ndvi_data$n_content)

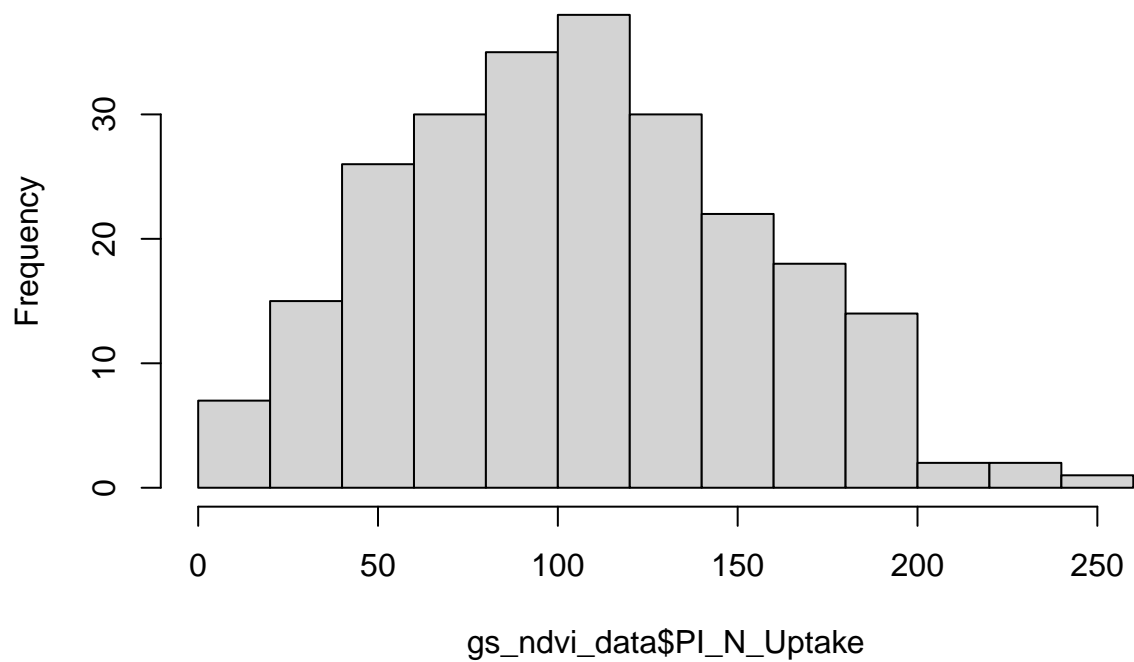
```

Histogram of gs_ndvi_data\$n_content



```
hist(gs_ndvi_data$PI_N_Uptake)
```

Histogram of gs_ndvi_data\$PI_N_Uptake



Yield Data

```
yield_data <- read_csv(file = "DATA/yield_data.csv")
```

```
## Rows: 672 Columns: 34
## -- Column specification -----
## Delimiter: ","
## chr (2): site_year, Received_TopDress
## dbl (32): year, Block, MainPlot, SubPlot, exp_plot_number, N_level, TopDress...
##
## 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.
```

```
str(yield_data , give.attr = FALSE)
```

```
## spec_tbl_df [672 x 34] (S3: spec_tbl_df/tbl_df/tbl/data.frame)
## $ site_year      : chr [1:672] "Davis-16" "Davis-16" "Davis-16" "Davis-16" ...
## $ year           : num [1:672] 2016 2016 2016 2016 2016 ...
## $ Block          : num [1:672] 1 1 1 1 1 2 2 2 2 2 ...
## $ MainPlot       : num [1:672] 1 2 3 4 5 1 2 3 4 5 ...
## $ SubPlot        : num [1:672] 3 1 3 2 1 2 3 2 3 2 ...
## $ exp_plot_number : num [1:672] 101 102 103 104 105 201 202 203 204 205 ...
## $ N_level        : num [1:672] 75 225 0 125 175 75 0 225 175 125 ...
## $ TopDress       : num [1:672] 0 0 0 0 0 0 0 0 0 0 ...
## $ SeasonalNRate  : num [1:672] 75 225 0 125 175 75 0 225 175 125 ...
## $ N_level_kgha   : num [1:672] 75 225 0 125 175 75 0 225 175 125 ...
## $ TopDress_kgha  : num [1:672] 0 0 0 0 0 0 0 0 0 0 ...
## $ SeasonalNRate_kgha : num [1:672] 75 225 0 125 175 75 0 225 175 125 ...
## $ Received_TopDress : chr [1:672] "NO" "NO" "NO" "NO" ...
## $ tare           : num [1:672] 1220 1220 1220 1220 1220 1220 1220 1220 1220 1220 ...
## $ FW1PlusTare    : num [1:672] 5662 5298 5256 4846 5220 ...
## $ FW2PlusTare    : num [1:672] 4610 5438 1220 4278 5156 ...
## $ SSFWPlusTare   : num [1:672] 3070 3262 2266 2482 2650 ...
## $ SSODW          : num [1:672] 692 705 519 541 538 ...
## $ HarvestBagPlusTie : num [1:672] 121 122 122 121 122 ...
## $ Grain1PlusPaperBag1 : num [1:672] 292 271 188 215 208 ...
## $ PaperBag1      : num [1:672] 7.92 7.92 7.92 7.92 7.92 7.92 7.92 7.92 7.92 7.92 ...
## $ Grain2PlusPaperBag2 : num [1:672] 7.01 7.01 7.01 7.01 7.01 7.01 7.01 7.01 7.01 7.01 ...
## $ PaperBag2      : num [1:672] 7.01 7.01 7.01 7.01 7.01 7.01 7.01 7.01 7.01 7.01 ...
## $ StrawSampleSize : num [1:672] 8.27 7.74 7.9 8.38 7.52 ...
## $ StrawN         : num [1:672] 55.1 55.2 49.4 47.1 45.3 ...
## $ GrainSampleSize : num [1:672] 4.95 4.97 5.04 5.02 4.57 ...
## $ GrainN         : num [1:672] 54.5 58.6 47.4 52.5 46.1 ...
## $ DaysPI2Harvest  : num [1:672] 102 102 102 102 102 102 102 102 102 102 ...
## $ SeedTray1.1     : num [1:672] 243 243 243 243 243 243 243 243 243 243 ...
## $ SeedTray1.2     : num [1:672] 0.62 0.62 0.62 0.64 0.64 0.62 0.62 0.62 0.62 0.62 ...
## $ Grain3PlusSeedTray1 : num [1:672] 435 514 426 455 447 ...
## $ MoistureContentGrain3 : num [1:672] 11.2 11.2 11.2 11.2 11.2 11.2 11.2 11.2 11.2 11.2 ...
## $ SeedTray2       : num [1:672] 244 244 244 244 244 ...
## $ Grain4PlusSeedTray2 : num [1:672] 254 254 254 254 254 ...
```

```
yield_data <- yield_data %>%
```

```
  filter(!year %in% c( "2016") & TopDress_kgha == 0 & N_level_kgha != 275) #removing the years and N r
```

```
yield_data <- yield_data %>%
```

```

mutate(
  site_year = factor(site_year),
  year = factor(year),
  Block = factor(Block),
  MainPlot = factor(MainPlot),
  exp_plot_number = factor(exp_plot_number),
  N_level = factor(N_level),
  SubPlot = factor(SubPlot),
  TopDress = factor(TopDress),
  SeasonalNRate_f = factor(SeasonalNRate),
  N_level_kgha_f = factor(N_level_kgha),
  TopDress_kgha_f = factor(TopDress_kgha),
  SeasonalNRate_kgha_f = factor(SeasonalNRate_kgha)
) #changes these columns to factor

str(yield_data , give.attr = FALSE)

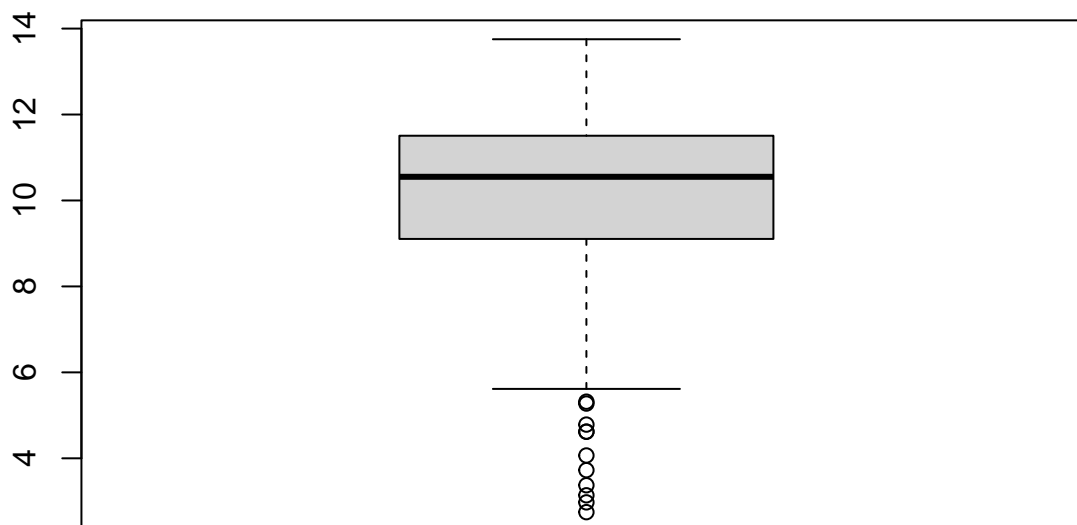
## tibble [240 x 38] (S3: tbl_df/tbl/data.frame)
## $ site_year      : Factor w/ 10 levels "Arbuckle-18",...: 7 7 7 7 7 7 7 7 7 7 ...
## $ year           : Factor w/ 3 levels "2017","2018",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ Block          : Factor w/ 4 levels "1","2","3","4": 1 1 1 1 1 1 2 2 2 2 ...
## $ MainPlot       : Factor w/ 7 levels "1","2","3","4",...: 1 2 3 4 6 7 2 3 4 5 ...
## $ SubPlot        : Factor w/ 3 levels "1","2","3": 2 3 1 1 1 2 1 1 2 3 ...
## $ exp_plot_number : Factor w/ 27 levels "101","102","103",...: 1 2 3 4 6 7 9 10 11 12 ...
## $ N_level        : Factor w/ 11 levels "0","45","75",...: 11 1 6 8 2 3 3 6 1 8 ...
## $ TopDress       : Factor w/ 1 level "0": 1 1 1 1 1 1 1 1 1 1 ...
## $ SeasonalNRate   : num [1:240] 225 0 125 175 45 75 75 125 0 175 ...
## $ N_level_kgha    : num [1:240] 225 0 125 175 45 75 75 125 0 175 ...
## $ TopDress_kgha   : num [1:240] 0 0 0 0 0 0 0 0 0 0 ...
## $ SeasonalNRate_kgha : num [1:240] 225 0 125 175 45 75 75 125 0 175 ...
## $ Received_TopDress : chr [1:240] "NO" "NO" "NO" "NO" ...
## $ tare           : num [1:240] 1220 1220 1220 1220 1220 1220 1220 1220 1220 1220 ...
## $ FW1PlusTare     : num [1:240] 4360 4818 5376 5598 4852 ...
## $ FW2PlusTare     : num [1:240] 3254 1220 1220 1220 1220 ...
## $ SSFWPlusTare    : num [1:240] 2324 1814 1994 2126 2106 ...
## $ SSODW           : num [1:240] 632 479 570 562 583 ...
## $ HarvestBagPlusTie : num [1:240] 162 176 169 121 120 ...
## $ Grain1PlusPaperBag1 : num [1:240] 230 173 211 233 265 ...
## $ PaperBag1       : num [1:240] 7.96 7.96 7.96 7.96 7.96 7.96 7.96 7.96 7.96 7.96 ...
## $ Grain2PlusPaperBag2 : num [1:240] 7.01 7.01 7.01 7.01 7.01 7.01 7.01 7.01 7.01 7.01 ...
## $ PaperBag2       : num [1:240] 7.01 7.01 7.01 7.01 7.01 7.01 7.01 7.01 7.01 7.01 ...
## $ StrawSampleSize : num [1:240] 8.25 7.96 7.76 7.89 8.45 ...
## $ StrawN          : num [1:240] 63.5 36 46.1 55.9 39.7 ...
## $ GrainSampleSize : num [1:240] 4.97 5.52 5.23 5.72 5.82 ...
## $ GrainN          : num [1:240] 60.5 50 56 66.7 51.1 ...
## $ DaysPI2Harvest   : num [1:240] 76 76 76 76 76 76 76 76 76 ...
## $ SeedTray1.1      : num [1:240] 243 243 243 243 243 243 243 243 243 ...
## $ SeedTray1.2      : num [1:240] 0.6 0.6 0.6 0.6 0.6 0.6 0.6 0.6 0.6 ...
## $ Grain3PlusSeedTray1 : num [1:240] 470 410 449 471 504 ...
## $ MoistureContentGrain3: num [1:240] 11.2 11.2 11.2 11.2 11.2 11.2 11.2 11.2 11.2 ...
## $ SeedTray2        : num [1:240] 244 244 244 244 244 ...
## $ Grain4PlusSeedTray2 : num [1:240] 254 254 254 254 254 ...
## $ SeasonalNRate_f   : Factor w/ 11 levels "0","45","75",...: 11 1 6 8 2 3 3 6 1 8 ...
## $ N_level_kgha_f    : Factor w/ 11 levels "0","45","75",...: 10 1 5 8 2 3 3 5 1 8 ...

```

```
## $ TopDress_kgha_f      : Factor w/ 1 level "0": 1 1 1 1 1 1 1 1 1 1 ...
## $ SeasonalNRate_kgha_f : Factor w/ 11 levels "0","45","75",...: 10 1 5 8 2 3 3 5 1 8 ...

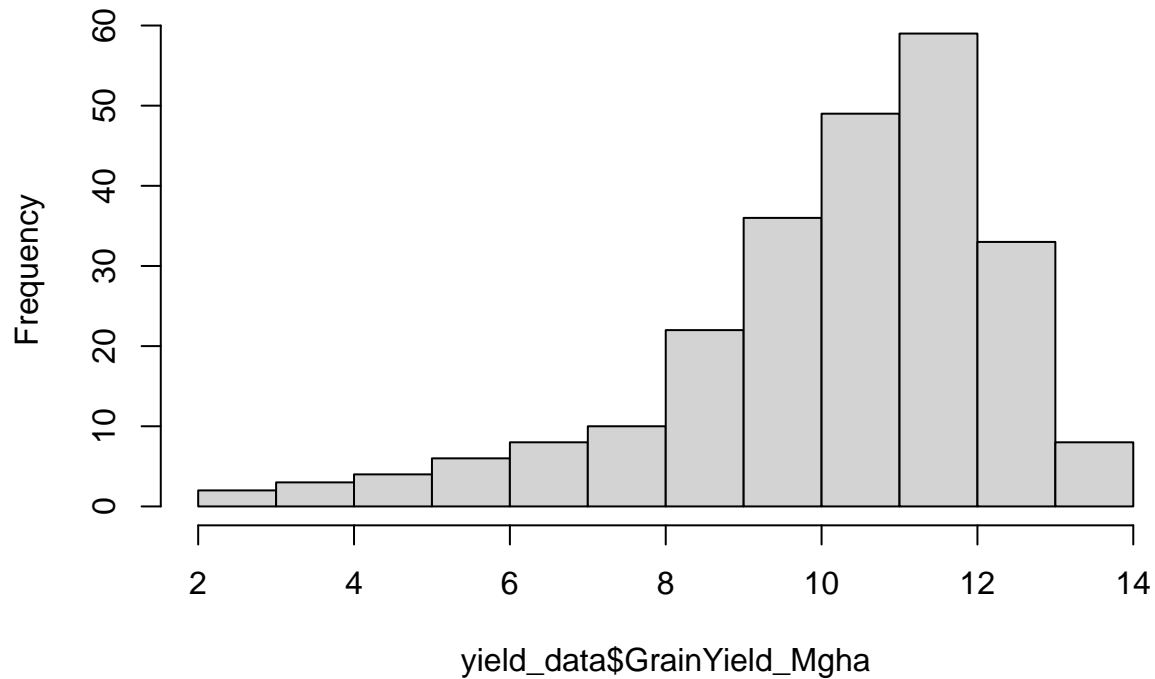
yield_data <- yield_data %>%
  mutate(
    FW1net = FW1PlusTare - tare,
    FW2net = FW2PlusTare - tare,
    TotalFW = FW1net + FW2net,
    SSFWnet = SSFWPlusTare - tare,
    Ratio = SSFWnet / TotalFW,
    SSODWnet = SSODW - HarvestBagPlusTie,
    SeedTray1 = SeedTray1.1 + SeedTray1.2, #adds the decimal to the 243g to get the tare weight for the
    Grain3net = Grain3PlusSeedTray1 - SeedTray1, #subtract tare of seed tray from grain3. Grain3 is the
    Grain4net = Grain4PlusSeedTray2 - SeedTray2, #grain4 is the amount of grain removed for ballmilling
    Grain2net = Grain2PlusPaperBag2 - PaperBag2, #yield component grain sample
    Grain2net = Grain2net * Ratio, #this essentially subsamples the yield component grain sample
    GrainNet = Grain3net + Grain4net + Grain2net, #add the grain removed for ball milling and yield comp
    GrainRing = GrainNet / Ratio, #the amount of grain in the entire m^2 ring in grams
    GrainYield = GrainRing * 10, #g/m^2 to kg/ha
    GrainYield_kgha = GrainYield * ((100-MoistureContentGrain3)/86), #corrects for 14% moisture based on
    GrainYield_Mgha = GrainYield_kgha / 1000, #converts kg/ha to Mg/ha
    Grain5 = GrainRing * ((100-MoistureContentGrain3)/98.1), #grain in the ring if the subsample was at
    Grain6 = GrainNet * ((100-MoistureContentGrain3)/98.1), #grain in the subsample if it was at 1.9% m
    StrawSS = SSODWnet - Grain6, #just straw in subsample in grams
    StrawRing = StrawSS / Ratio, #straw in ring in grams i.e g/m2
    StrawNcon = StrawN / StrawSampleSize,
    StrawNup = (StrawRing * StrawNcon) / 100, #straw Nup divide by 100 to convert mg/m2 to kg/ha - this
    GrainNcon = (GrainN / GrainSampleSize), #grain in ring in kg/ha
    GrainNup = (Grain5 * GrainNcon) / 100, #grain Nup divide by 100 to convert mg N/m2 to kg N/ha
    TotalSeasonalNup = StrawNup + GrainNup, #in kg/ha
    HarvestIndex = Grain5 / (Grain5 + StrawRing),
    Moisture = SSFWnet / SSODWnet
  )

boxplot(yield_data$GrainYield_Mgha)
```

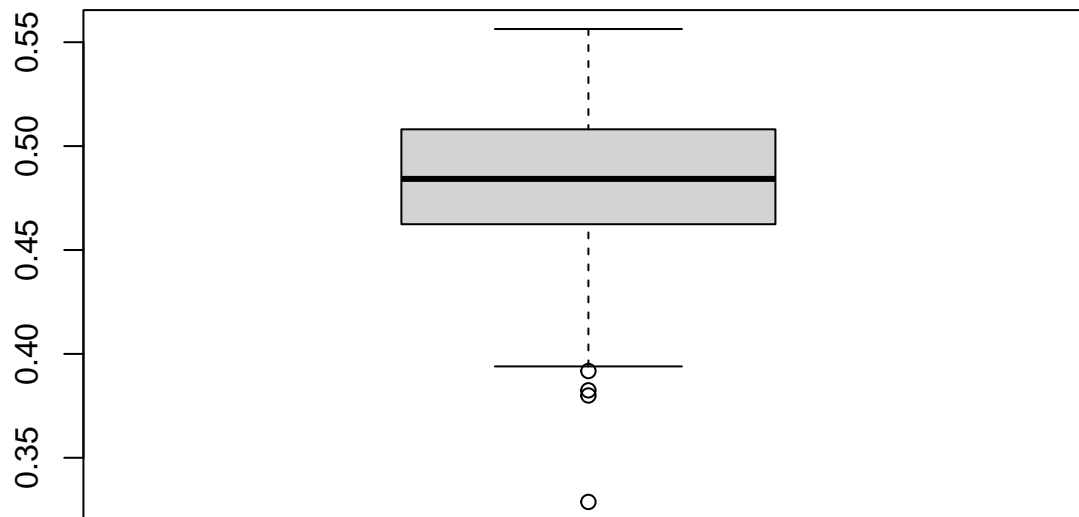



```
hist(yield_data$GrainYield_Mgha)
```

Histogram of yield_data\$GrainYield_Mgha

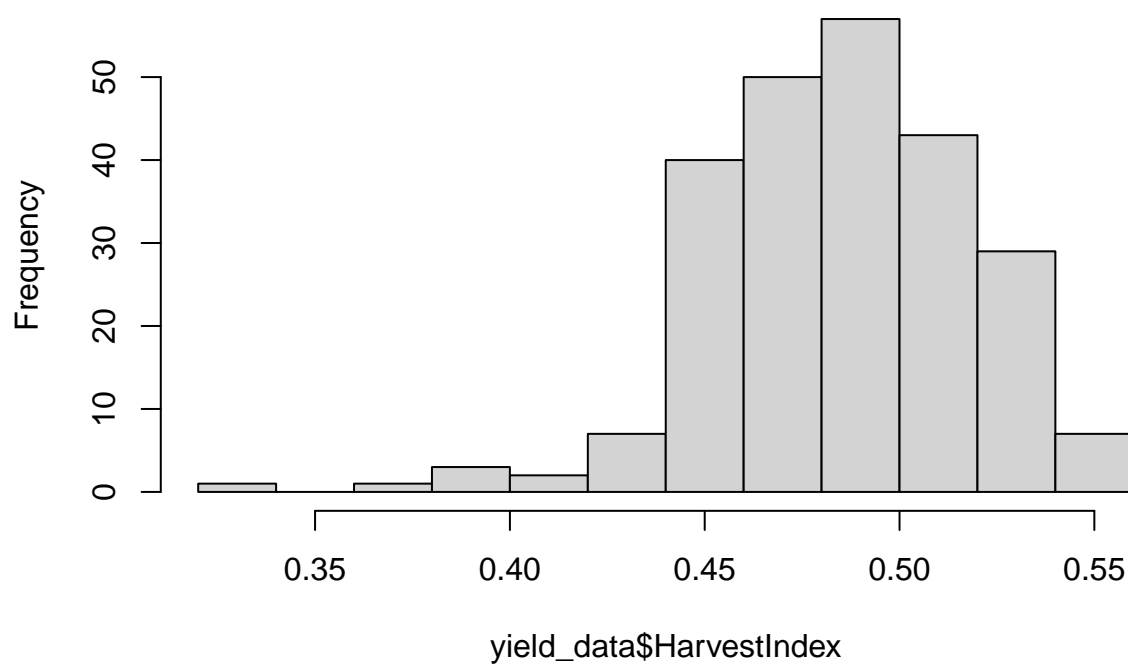


```
boxplot(yield_data$HarvestIndex)
```

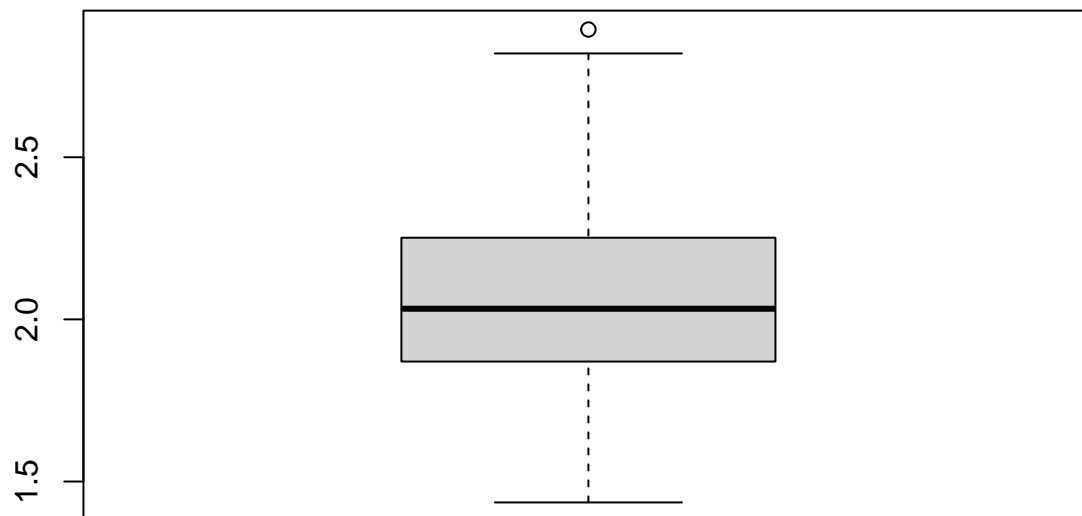


```
hist(yield_data$HarvestIndex)
```

Histogram of yield_data\$HarvestIndex

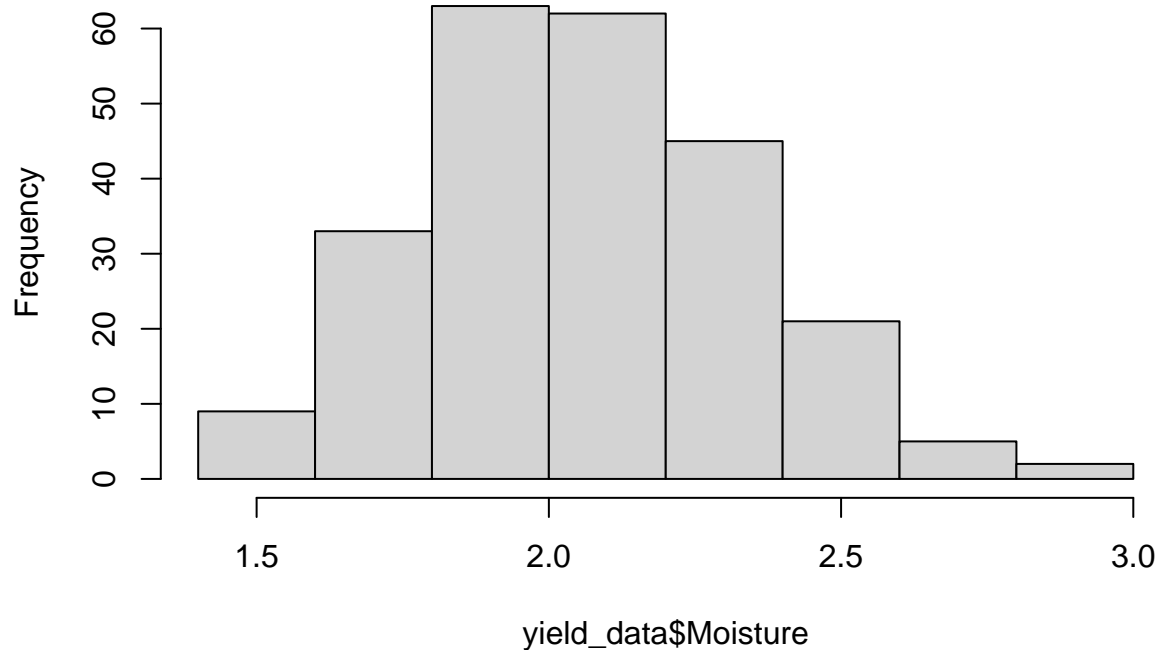


```
boxplot(yield_data$Moisture)
```



```
hist(yield_data$Moisture)
```

Histogram of yield_data\$Moisture



#the data looks good - don't see any unusual values

```
yield_data <- dplyr::select(yield_data,
  site_year,
  year,
  Block,
  MainPlot,
  SubPlot,
  exp_plot_number,
  N_level_kgha,
  N_level_kgha_f,
  TopDress_kgha,
  TopDress_kgha_f,
  GrainYield_Mgha
)
```

```
gs_data <- full_join(gs_ndvi_data , yield_data)
```

```
## Joining, by = c("site_year", "year", "exp_plot_number", "N_level_kgha")
```

```
gs_data <- dplyr::select(gs_data,
  site_year,
  year,
  exp_plot_number,
  Block,
  MainPlot,
  SubPlot,
  N_level_kgha,
```

```

      N_level_kgha_f,
      TopDress_kgha,
      TopDress_kgha_f ,
      PI_N_Uptake,
      NDVI,
      GrainYield_Mgha) #reorders the columns

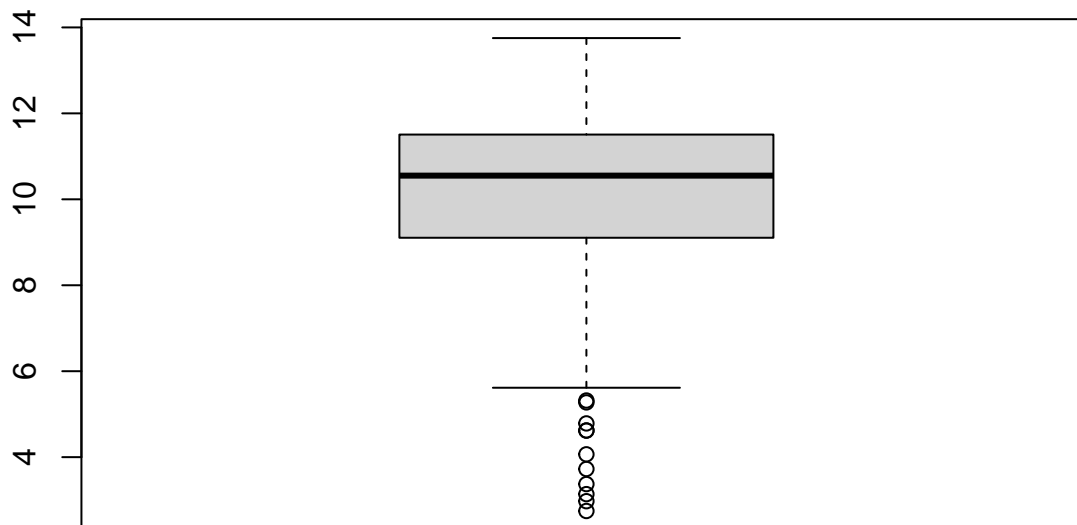
gs_data$site_year <- as.factor(gs_data$site_year)

str(gs_data , give.attr = FALSE)

## rowwise_df [240 x 13] (S3: rowwise_df/tbl_df/tbl/data.frame)
## $ site_year      : Factor w/ 10 levels "Nicolaus-17",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ year           : Factor w/ 3 levels "2017","2018",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ exp_plot_number: Factor w/ 27 levels "101","102","103",...: 1 2 3 4 6 7 9 10 11 12 ...
## $ Block          : Factor w/ 4 levels "1","2","3","4": 1 1 1 1 1 1 2 2 2 2 ...
## $ MainPlot       : Factor w/ 7 levels "1","2","3","4",...: 1 2 3 4 6 7 2 3 4 5 ...
## $ SubPlot        : Factor w/ 3 levels "1","2","3": 2 3 1 1 1 2 1 1 2 3 ...
## $ N_level_kgha   : num [1:240] 225 0 125 175 45 75 75 125 0 175 ...
## $ N_level_kgha_f : Factor w/ 11 levels "0","45","75",...: 10 1 5 8 2 3 3 5 1 8 ...
## $ TopDress_kgha  : num [1:240] 0 0 0 0 0 0 0 0 0 0 ...
## $ TopDress_kgha_f: Factor w/ 1 level "0": 1 1 1 1 1 1 1 1 1 1 ...
## $ PI_N_Uptake    : num [1:240] 158 78 142 197 95 ...
## $ NDVI           : num [1:240] 0.777 0.493 0.683 0.787 0.683 ...
## $ GrainYield_Mgha: num [1:240] 11.4 11 12 11.9 11.5 ...

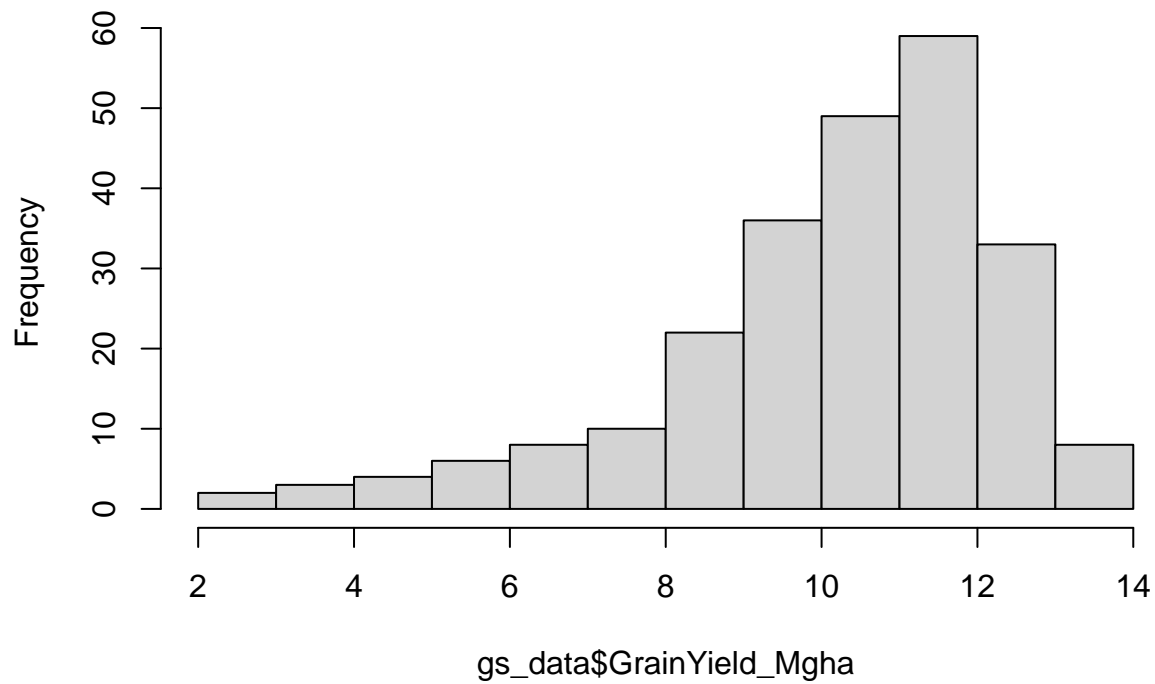
```

```
boxplot(gs_data$GrainYield_Mgha)
```



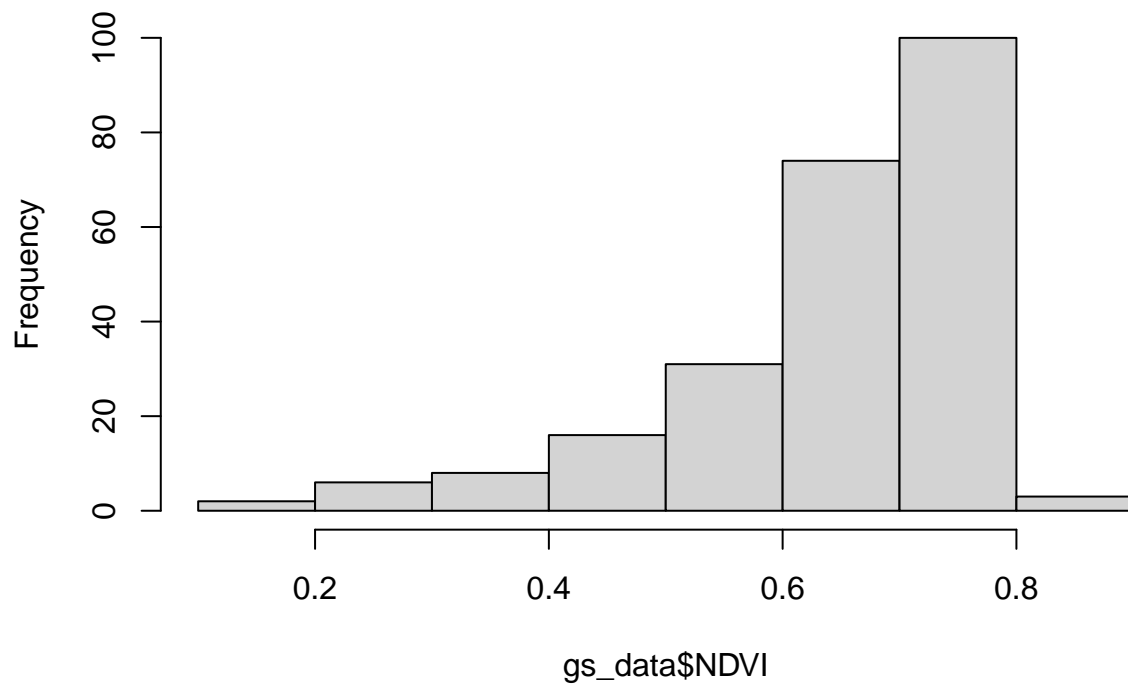
```
hist(gs_data$GrainYield_Mgha)
```

Histogram of gs_data\$GrainYield_Mgha

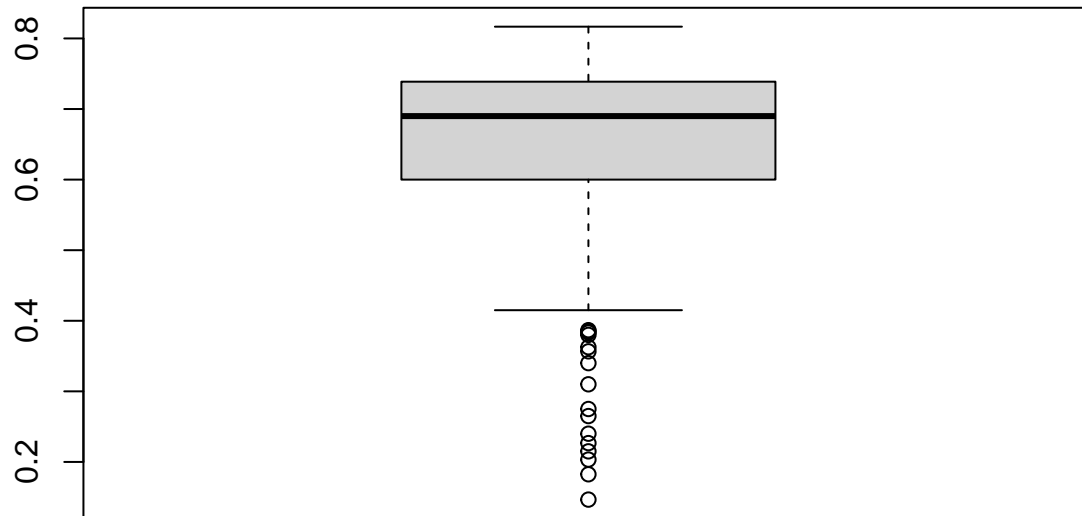


```
hist(gs_data$NDVI)
```

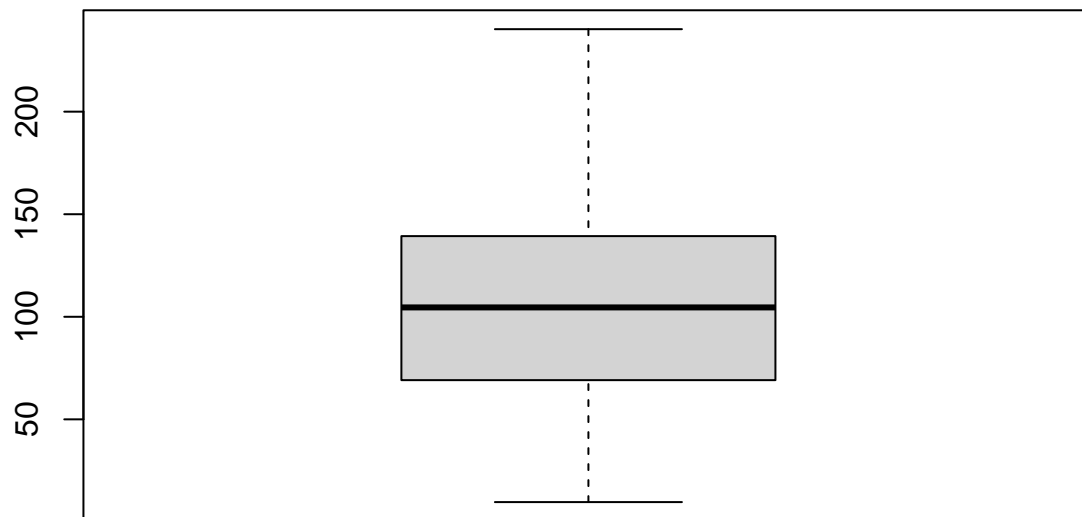
Histogram of gs_data\$NDVI



```
boxplot(gs_data$NDVI)
```

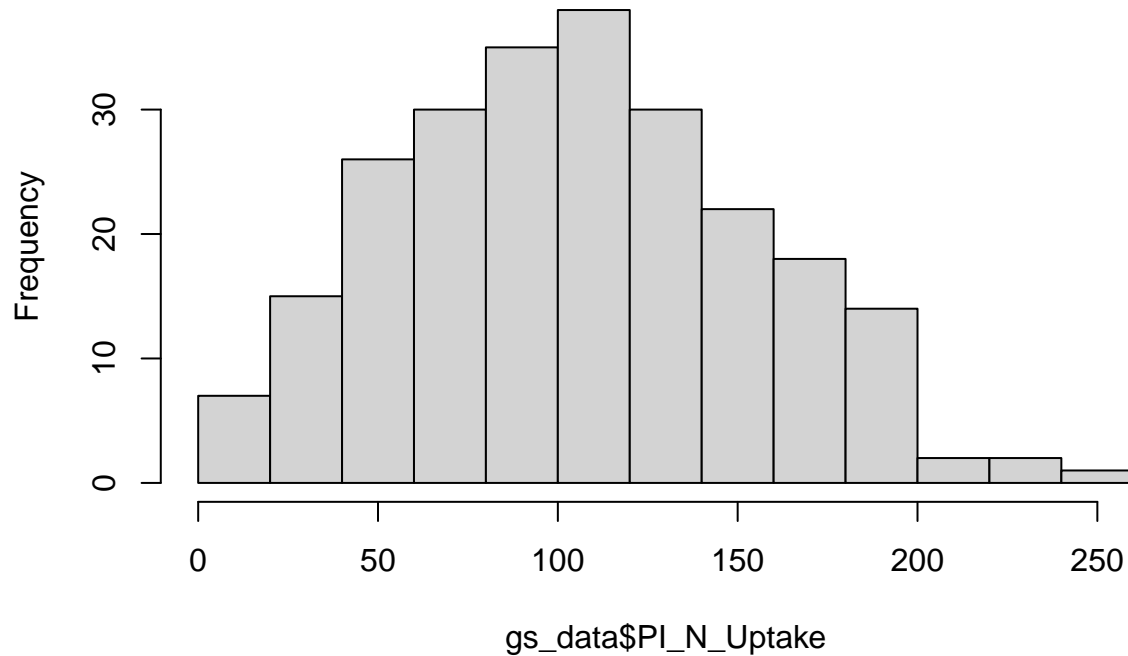


```
boxplot(gs_data$PI_N_Uptake)
```



```
hist(gs_data$PI_N_Uptake)
```

Histogram of gs_data\$PI_N_Uptake



#Overall data looks good -- no errors of data entry

Calculating GS RI

```
max_gs_ndvi <- gs_data %>%
  filter(N_level_kgha_f %in% c(225, 235)) %>%
  dplyr::select(site_year , NDVI) %>%
  group_by(site_year) %>%
  summarise(mean_NDVI = mean(NDVI))

nic17 <- subset(max_gs_ndvi, site_year == "Nicolaus-17")
nic17maxNDVI <- nic17$mean_NDVI
nic17maxNDVI <- as.numeric(nic17maxNDVI)
nic17maxNDVI
```

```
## [1] 0.7858333
```

```
wil17 <- subset(max_gs_ndvi, site_year == "Williams-17")
wil17maxNDVI <- wil17$mean_NDVI
wil17maxNDVI <- as.numeric(wil17maxNDVI)
wil17maxNDVI
```

```
## [1] 0.7925
```

```
arb18 <- subset(max_gs_ndvi, site_year == "Arbuckle-18")
arb18maxNDVI <- arb18$mean_NDVI
arb18maxNDVI <- as.numeric(arb18maxNDVI)
arb18maxNDVI
```

```
## [1] 0.73875
```

```

biggs18 <- subset(max_gs_ndvi, site_year == "Biggs-18")
biggs18maxNDVI <- biggs18$mean_NDVI
biggs18maxNDVI <- as.numeric(biggs18maxNDVI)
biggs18maxNDVI

## [1] 0.784375

mry18 <- subset(max_gs_ndvi, site_year == "Marysville-18")
mry18maxNDVI <- mry18$mean_NDVI
mry18maxNDVI <- as.numeric(mry18maxNDVI)
mry18maxNDVI

## [1] 0.728125

nic18 <- subset(max_gs_ndvi, site_year == "Nicolaus-18")
nic18maxNDVI <- nic18$mean_NDVI
nic18maxNDVI <- as.numeric(nic18maxNDVI)
nic18maxNDVI

## [1] 0.765

arb19 <- subset(max_gs_ndvi, site_year == "Arbuckle-19")
arb19maxNDVI <- arb19$mean_NDVI
arb19maxNDVI <- as.numeric(arb19maxNDVI)
arb19maxNDVI

## [1] 0.72

davis19 <- subset(max_gs_ndvi, site_year == "Davis-19")
davis19maxNDVI <- davis19$mean_NDVI
davis19maxNDVI <- as.numeric(davis19maxNDVI)
davis19maxNDVI

## [1] 0.613125

mry19 <- subset(max_gs_ndvi, site_year == "Marysville-19")
mry19maxNDVI <- mry19$mean_NDVI
mry19maxNDVI <- as.numeric(mry19maxNDVI)
mry19maxNDVI

## [1] 0.745625

res19 <- subset(max_gs_ndvi, site_year == "RES-19")
res19maxNDVI <- res19$mean_NDVI
res19maxNDVI <- as.numeric(res19maxNDVI)
res19maxNDVI

## [1] 0.66625

gs_data <- gs_data %>%
  mutate(max_NDVI = case_when(
    site_year == "Nicolaus-17" ~ nic17maxNDVI ,
    site_year == "Williams-17" ~ wil17maxNDVI ,
    site_year == "Arbuckle-18" ~ arb18maxNDVI ,
    site_year == "Biggs-18" ~ biggs18maxNDVI ,
    site_year == "Marysville-18" ~ mry18maxNDVI ,
    site_year == "Nicolaus-18" ~ nic18maxNDVI ,
    site_year == "Arbuckle-19" ~ arb19maxNDVI ,
    site_year == "Davis-19" ~ davis19maxNDVI ,

```



```

        site_year == "Marysville-19" ~ mry19maxNDVI ,
        site_year == "RES-19" ~ res19maxNDVI)
)

gs_data <- gs_data %>%
  mutate(gs_NDVI_Response_Index = case_when(
    site_year == "Nicolaus-17" ~ nic17maxNDVI / NDVI,
    site_year == "Williams-17" ~ wil17maxNDVI / NDVI,
    site_year == "Arbuckle-18" ~ arb18maxNDVI / NDVI ,
    site_year == "Biggs-18" ~ biggs18maxNDVI / NDVI ,
    site_year == "Marysville-18" ~ mry18maxNDVI / NDVI ,
    site_year == "Nicolaus-18" ~ nic18maxNDVI / NDVI,
    site_year == "Arbuckle-19" ~ arb19maxNDVI / NDVI,
    site_year == "Davis-19" ~ davis19maxNDVI / NDVI,
    site_year == "Marysville-19" ~ mry19maxNDVI / NDVI,
    site_year == "RES-19" ~ res19maxNDVI / NDVI
  )) #calculates NDVI response index

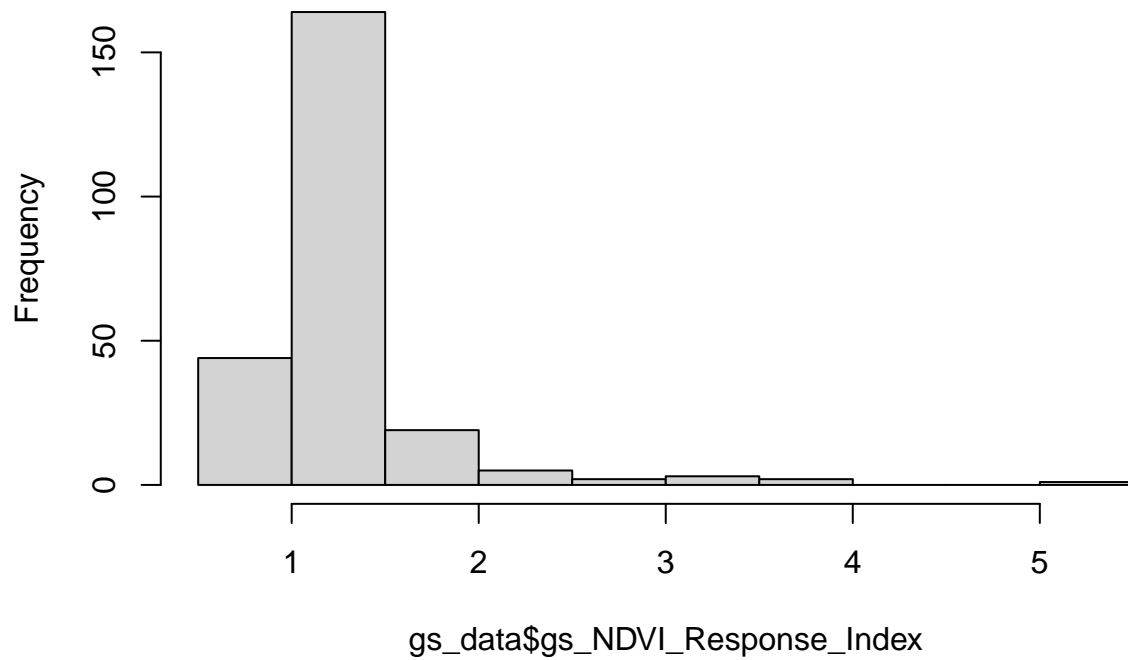
str(gs_data , give.attr = FALSE)

## rowwise_df [240 x 15] (S3: rowwise_df/tbl_df/tbl/data.frame)
## $ site_year      : Factor w/ 10 levels "Nicolaus-17",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ year           : Factor w/ 3 levels "2017","2018",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ exp_plot_number : Factor w/ 27 levels "101","102","103",...: 1 2 3 4 6 7 9 10 11 12 ...
## $ Block          : Factor w/ 4 levels "1","2","3","4": 1 1 1 1 1 1 2 2 2 2 ...
## $ MainPlot        : Factor w/ 7 levels "1","2","3","4",...: 1 2 3 4 6 7 2 3 4 5 ...
## $ SubPlot         : Factor w/ 3 levels "1","2","3": 2 3 1 1 1 2 1 1 2 3 ...
## $ N_level_kgha     : num [1:240] 225 0 125 175 45 75 75 125 0 175 ...
## $ N_level_kgha_f   : Factor w/ 11 levels "0","45","75",...: 10 1 5 8 2 3 3 5 1 8 ...
## $ TopDress_kgha    : num [1:240] 0 0 0 0 0 0 0 0 0 0 ...
## $ TopDress_kgha_f  : Factor w/ 1 level "0": 1 1 1 1 1 1 1 1 1 1 ...
## $ PI_N_Uptake      : num [1:240] 158 78 142 197 95 ...
## $ NDVI             : num [1:240] 0.777 0.493 0.683 0.787 0.683 ...
## $ GrainYield_Mgha  : num [1:240] 11.4 11 12 11.9 11.5 ...
## $ max_NDVI         : num [1:240] 0.786 0.786 0.786 0.786 0.786 ...
## $ gs_NDVI_Response_Index: num [1:240] 1.012 1.593 1.15 0.999 1.15 ...

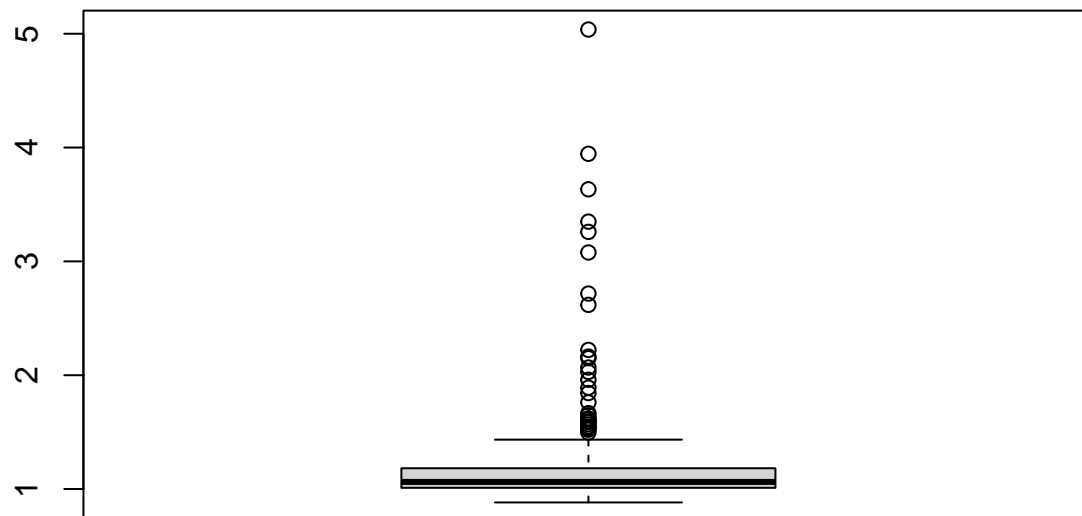
hist(gs_data$gs_NDVI_Response_Index)

```

Histogram of gs_data\$gs_NDVI_Response_Index

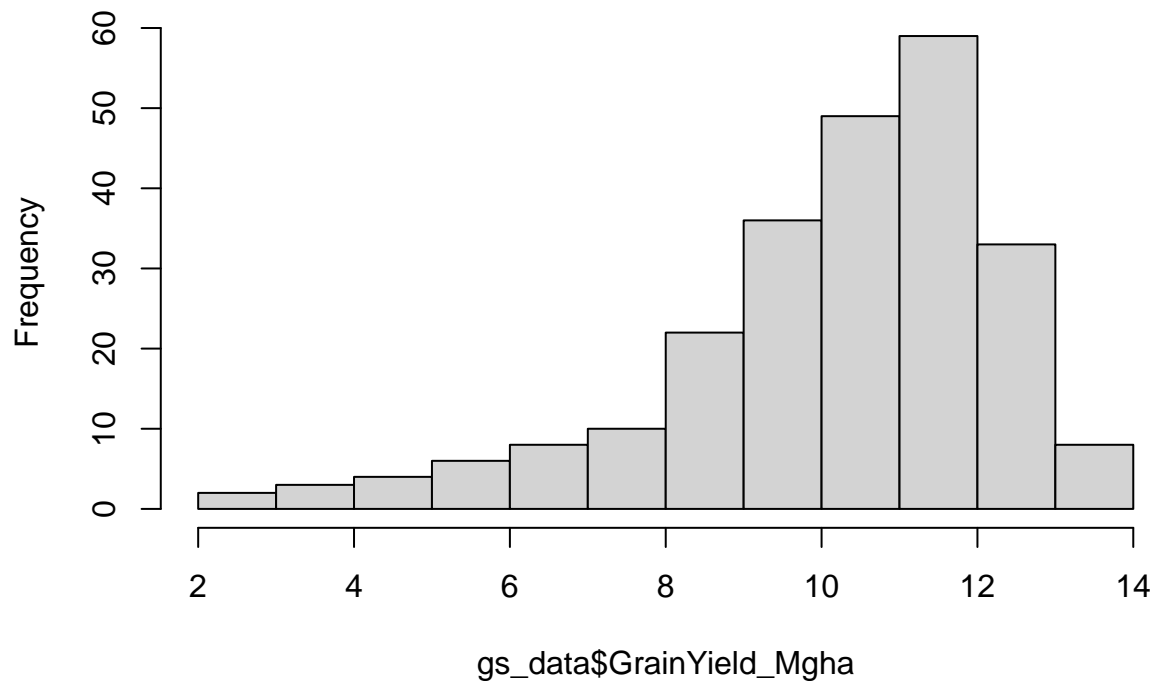


```
boxplot(gs_data$gs_NDVI_Response_Index)
```

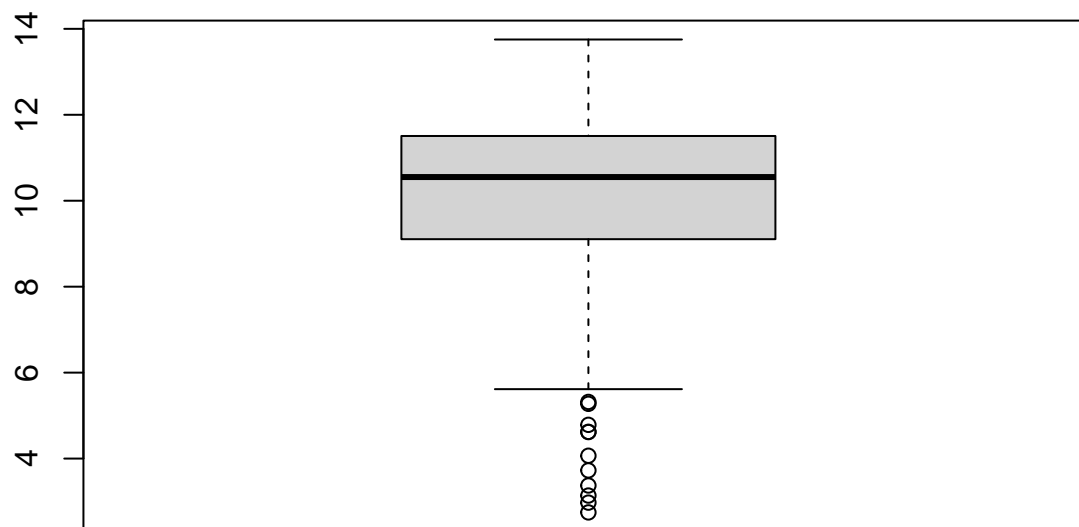


```
hist(gs_data$GrainYield_Mgha)
```

Histogram of gs_data\$GrainYield_Mgha



```
boxplot(gs_data$GrainYield_Mgha)
```



Drone Data

```
drone_data <- read_csv(file = "DATA/PI_drone_data.csv")
```

```
## Rows: 248 Columns: 38
## -- Column specification -----
## Delimiter: ","
## chr (1): site_year
## dbl (37): year, exp_plot_number, Block, MainPlot, N_level, N_level_kgha, bio...
##
```

```
## 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.
```

```
str(drone_data , give.attr = FALSE)
```

```
## spec_tbl_df [248 x 38] (S3: spec_tbl_df/tbl_df/tbl/data.frame)
## $ site_year      : chr [1:248] "Nicolaus-17" "Nicolaus-17" "Nicolaus-17" "Nicolaus-17" ...
## $ year           : num [1:248] 2017 2017 2017 2017 2017 2017 ...
## $ exp_plot_number : num [1:248] 101 102 103 104 105 106 107 201 202 203 ...
## $ Block          : num [1:248] 1 1 1 1 1 1 2 2 2 ...
## $ MainPlot       : num [1:248] 1 2 3 4 5 6 7 1 2 3 ...
## $ N_level        : num [1:248] 225 0 125 175 275 45 75 275 75 125 ...
## $ N_level_kgha    : num [1:248] 225 0 125 175 275 45 75 275 75 125 ...
## $ biomass_plus_bag_g : num [1:248] 361 264 318 360 394 ...
## $ ring_size      : num [1:248] 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 ...
## $ paper_bag_g     : num [1:248] 45.5 45.5 45.5 45.5 45.5 45.5 45.5 45.5 45.5 45.5 ...
## $ num_of_paper_bags : num [1:248] 1 1 1 1 1 1 1 1 1 1 ...
## $ sample_weight_mg : num [1:248] 3.17 3.42 3.09 3.07 3.35 ...
## $ sample_N_uq     : num [1:248] 79.4 61.1 80.7 95.9 111.3 ...
## $ greenmean       : num [1:248] 0.0467 0.0581 0.0498 0.0488 0.0505 0.0608 0.0589 0.0461 0.0537 0.0...
## $ greenmedia      : num [1:248] NA NA NA NA NA NA NA NA NA NA ...
## $ greenstdev      : num [1:248] NA NA NA NA NA NA NA NA NA NA ...
## $ greenmin        : num [1:248] NA NA NA NA NA NA NA NA NA NA ...
## $ greenmax        : num [1:248] NA NA NA NA NA NA NA NA NA NA ...
## $ bluemean        : num [1:248] NA NA NA NA NA NA NA NA NA NA ...
## $ bluedmedian     : num [1:248] NA NA NA NA NA NA NA NA NA NA ...
## $ bluestdev       : num [1:248] NA NA NA NA NA NA NA NA NA NA ...
## $ bluemin         : num [1:248] NA NA NA NA NA NA NA NA NA NA ...
## $ bluemax         : num [1:248] NA NA NA NA NA NA NA NA NA NA ...
## $ redmean         : num [1:248] 0.023 0.0286 0.0243 0.0237 0.0245 0.0292 0.0281 0.0234 0.0263 0.0...
## $ redmedian       : num [1:248] NA NA NA NA NA NA NA NA NA NA ...
## $ redstdev        : num [1:248] NA NA NA NA NA NA NA NA NA NA ...
## $ redmin          : num [1:248] NA NA NA NA NA NA NA NA NA NA ...
## $ redmax          : num [1:248] NA NA NA NA NA NA NA NA NA NA ...
## $ edgemean        : num [1:248] 0.0738 0.0867 0.0768 0.0764 0.0791 0.0903 0.0877 0.0744 0.0812 0.0...
## $ edgedmedian     : num [1:248] NA NA NA NA NA NA NA NA NA NA ...
## $ edgestdev       : num [1:248] NA NA NA NA NA NA NA NA NA NA ...
## $ edgemin         : num [1:248] NA NA NA NA NA NA NA NA NA NA ...
## $ edgemax         : num [1:248] NA NA NA NA NA NA NA NA NA NA ...
## $ nirmean         : num [1:248] 0.55 0.357 0.482 0.537 0.599 ...
## $ nirmedian       : num [1:248] NA NA NA NA NA NA NA NA NA NA ...
## $ nirstdev        : num [1:248] NA NA NA NA NA NA NA NA NA NA ...
## $ nirmin          : num [1:248] NA NA NA NA NA NA NA NA NA NA ...
## $ nirmax          : num [1:248] NA NA NA NA NA NA NA NA NA NA ...
```

```
drone_data <- drone_data %>%
  filter(N_level_kgha != 275) %>%
  mutate(year = factor(year) ,
         exp_plot_number = factor(exp_plot_number) ,
         Block = factor(Block) ,
         MainPlot = factor(MainPlot) ,
         N_level = factor(N_level) ,
         N_level_kgha_f = factor(N_level_kgha)
  )
```

```
drone_data$site_year <- factor(drone_data$site_year , levels = c("Nicolaus-17" , "Williams-17" , "Arbuc
str(drone_data , give.attr = FALSE)
```

```
## tibble [240 x 39] (S3: tbl_df/tbl/data.frame)
## $ site_year      : Factor w/ 10 levels "Nicolaus-17",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ year           : Factor w/ 3 levels "2017","2018",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ exp_plot_number : Factor w/ 27 levels "101","102","103",...: 1 2 3 4 6 7 9 10 11 12 ...
## $ Block          : Factor w/ 4 levels "1","2","3","4": 1 1 1 1 1 1 2 2 2 2 ...
## $ MainPlot        : Factor w/ 7 levels "1","2","3","4",...: 1 2 3 4 6 7 2 3 4 5 ...
## $ N_level         : Factor w/ 11 levels "0","45","75",...: 11 1 6 8 2 3 3 6 1 8 ...
## $ N_level_kgha    : num [1:240] 225 0 125 175 45 75 75 125 0 175 ...
## $ biomass_plus_bag_g: num [1:240] 361 264 318 360 285 ...
## $ ring_size       : num [1:240] 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 ...
## $ paper_bag_g     : num [1:240] 45.5 45.5 45.5 45.5 45.5 45.5 45.5 45.5 45.5 45.5 ...
## $ num_of_paper_bags : num [1:240] 1 1 1 1 1 1 1 1 1 1 ...
## $ sample_weight_mg : num [1:240] 3.17 3.42 3.09 3.07 3.2 ...
## $ sample_N_ug     : num [1:240] 79.4 61.1 80.7 95.9 63.6 ...
## $ greenmean       : num [1:240] 0.0467 0.0581 0.0498 0.0488 0.0608 0.0589 0.0537 0.0488 0.0598 0.0
## $ greenmedia      : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ greenstdev      : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ greenmin        : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ greenmax        : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ bluemean        : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ bluedmedian     : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ bluestdev       : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ bluemin         : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ bluemax         : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ redmean         : num [1:240] 0.023 0.0286 0.0243 0.0237 0.0292 0.0281 0.0263 0.0237 0.0293 0.0
## $ redmedian       : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ redstdev        : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ redmin          : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ redmax          : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ edgemean        : num [1:240] 0.0738 0.0867 0.0768 0.0764 0.0903 0.0877 0.0812 0.0754 0.0873 0.0
## $ edgemedian      : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ edgestdev       : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ edgemin         : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ edgemax         : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ nirmean         : num [1:240] 0.55 0.357 0.482 0.537 0.431 ...
## $ nirmedian       : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ nirstdev        : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ nirmin          : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ nirmax          : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ N_level_kgha_f   : Factor w/ 11 levels "0","45","75",...: 10 1 5 8 2 3 3 5 1 8 ...
```

```
drone_data <- dplyr::select(drone_data ,
                             site_year,
                             year,
                             exp_plot_number,
                             Block,
                             MainPlot,
                             N_level,
                             N_level_kgha,
                             N_level_kgha_f,
```

```

        biomass_plus_bag_g,
        ring_size,
        paper_bag_g,
        num_of_paper_bags,
        sample_weight_mg,
        sample_N_ug,
        bluemean,
        greenmean,
        redmean,
        edgemean,
        nirmean
    )#selects the relevant columns

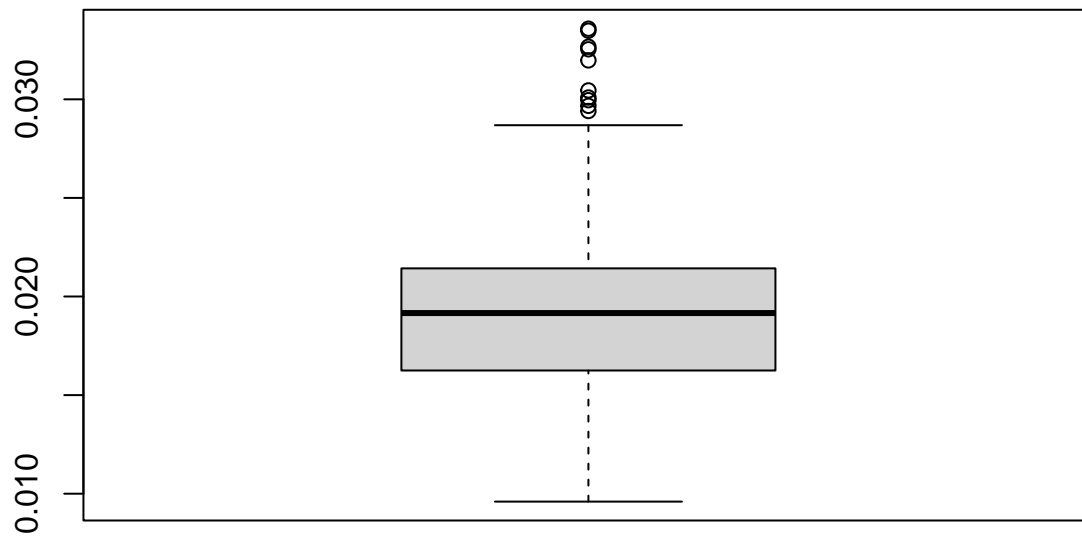
str(drone_data , give.attr = FALSE)

## tibble [240 x 19] (S3: tbl_df/tbl/data.frame)
## $ site_year      : Factor w/ 10 levels "Nicolaus-17",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ year           : Factor w/ 3 levels "2017","2018",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ exp_plot_number : Factor w/ 27 levels "101","102","103",...: 1 2 3 4 6 7 9 10 11 12 ...
## $ Block          : Factor w/ 4 levels "1","2","3","4": 1 1 1 1 1 1 2 2 2 2 ...
## $ MainPlot        : Factor w/ 7 levels "1","2","3","4",...: 1 2 3 4 6 7 2 3 4 5 ...
## $ N_level         : Factor w/ 11 levels "0","45","75",...: 11 1 6 8 2 3 3 6 1 8 ...
## $ N_level_kgha    : num [1:240] 225 0 125 175 45 75 75 125 0 175 ...
## $ N_level_kgha_f   : Factor w/ 11 levels "0","45","75",...: 10 1 5 8 2 3 3 5 1 8 ...
## $ biomass_plus_bag_g: num [1:240] 361 264 318 360 285 ...
## $ ring_size        : num [1:240] 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 ...
## $ paper_bag_g      : num [1:240] 45.5 45.5 45.5 45.5 45.5 45.5 45.5 45.5 45.5 45.5 ...
## $ num_of_paper_bags : num [1:240] 1 1 1 1 1 1 1 1 1 1 ...
## $ sample_weight_mg : num [1:240] 3.17 3.42 3.09 3.07 3.2 ...
## $ sample_N_ug       : num [1:240] 79.4 61.1 80.7 95.9 63.6 ...
## $ bluemean         : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ greenmean         : num [1:240] 0.0467 0.0581 0.0498 0.0488 0.0608 0.0589 0.0537 0.0488 0.0598 0.0...
## $ redmean           : num [1:240] 0.023 0.0286 0.0243 0.0237 0.0292 0.0281 0.0263 0.0237 0.0293 0.0...
## $ edgemean          : num [1:240] 0.0738 0.0867 0.0768 0.0764 0.0903 0.0877 0.0812 0.0754 0.0873 0.0...
## $ nirmean           : num [1:240] 0.55 0.357 0.482 0.537 0.431 ...

#visualize drone_data to look for outliers

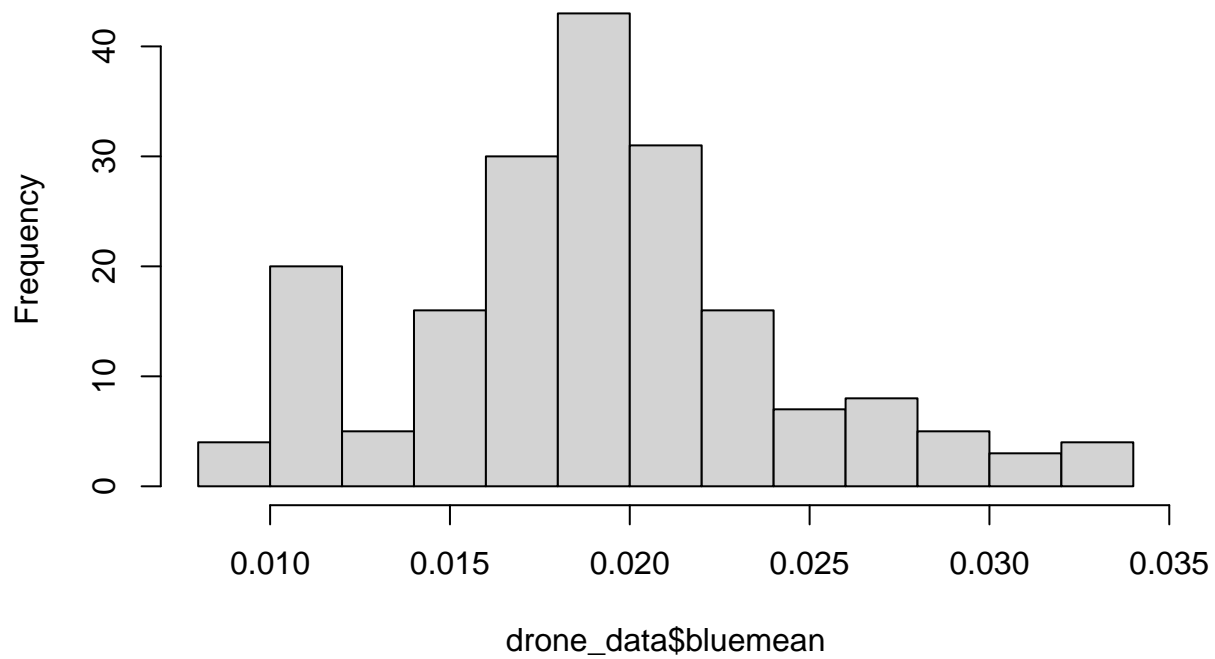
boxplot(drone_data$bluemean)

```

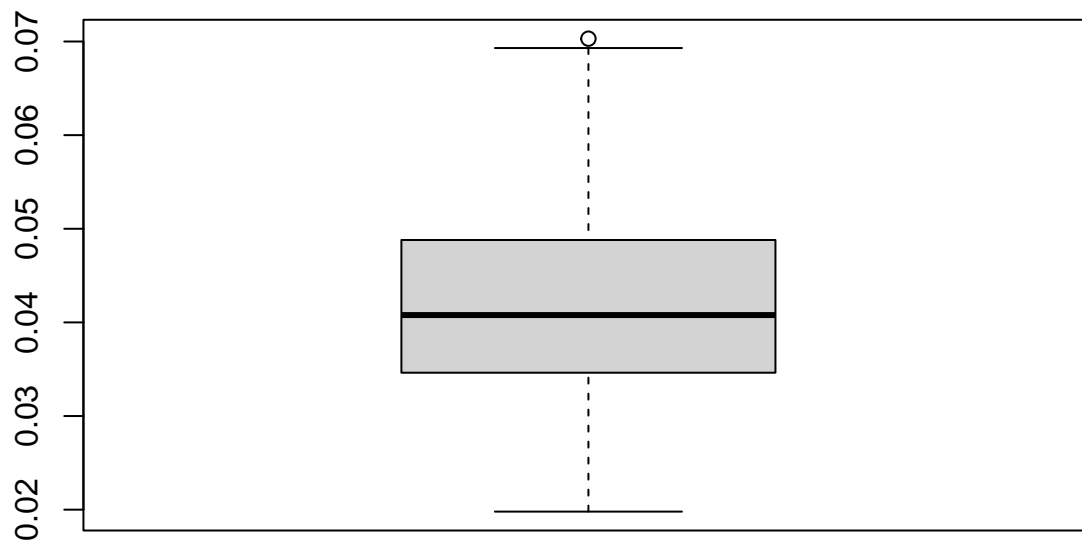


```
hist(drone_data$bluemean)
```

Histogram of drone_data\$bluemean

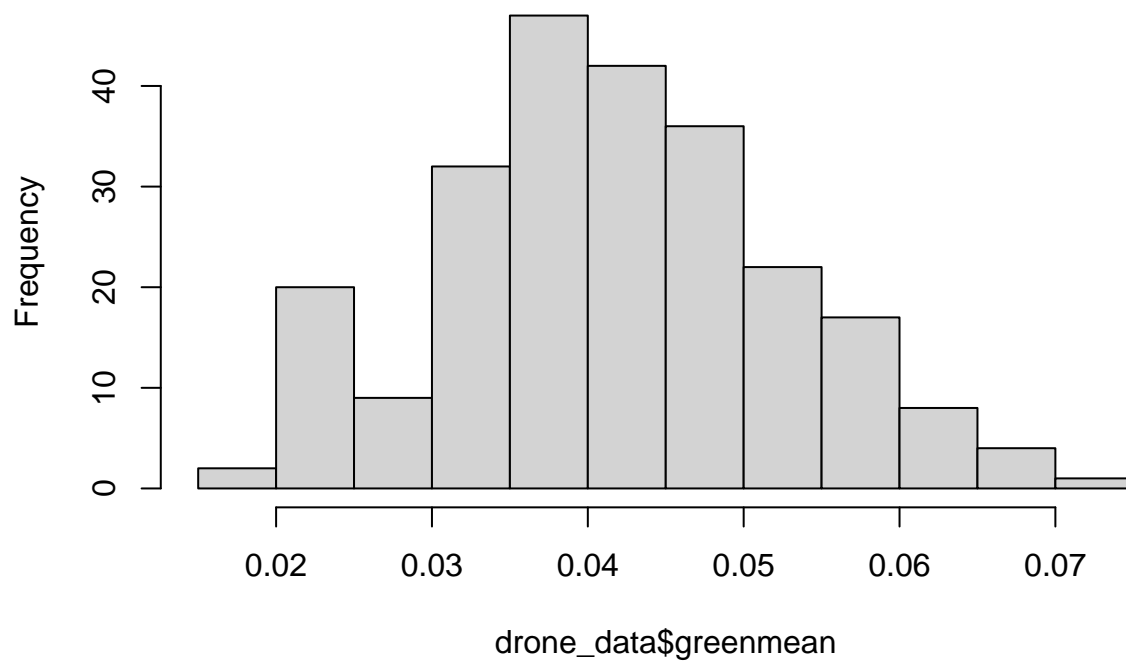


```
boxplot(drone_data$greenmean)
```

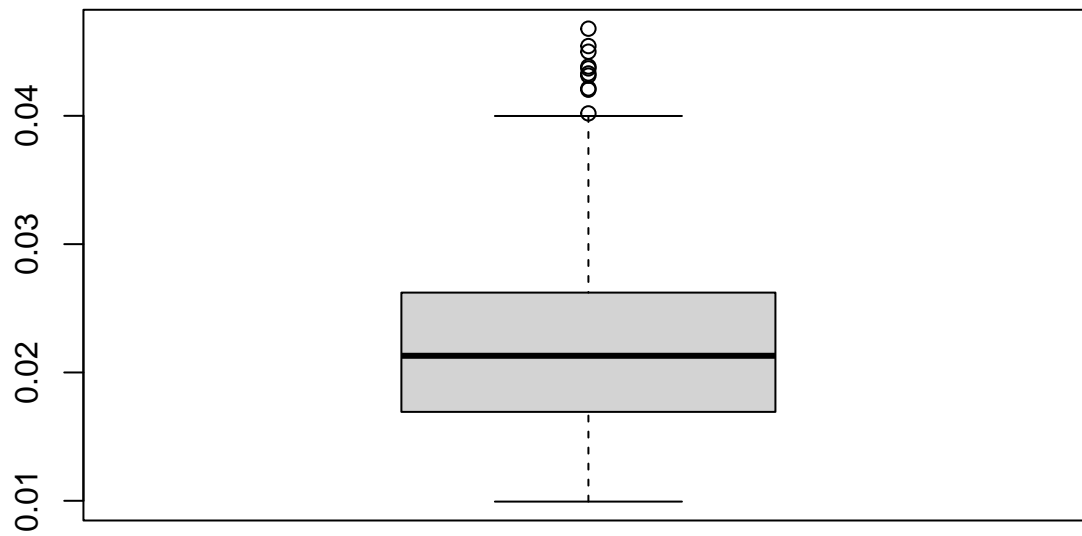


```
hist(drone_data$greenmean)
```

Histogram of drone_data\$greenmean

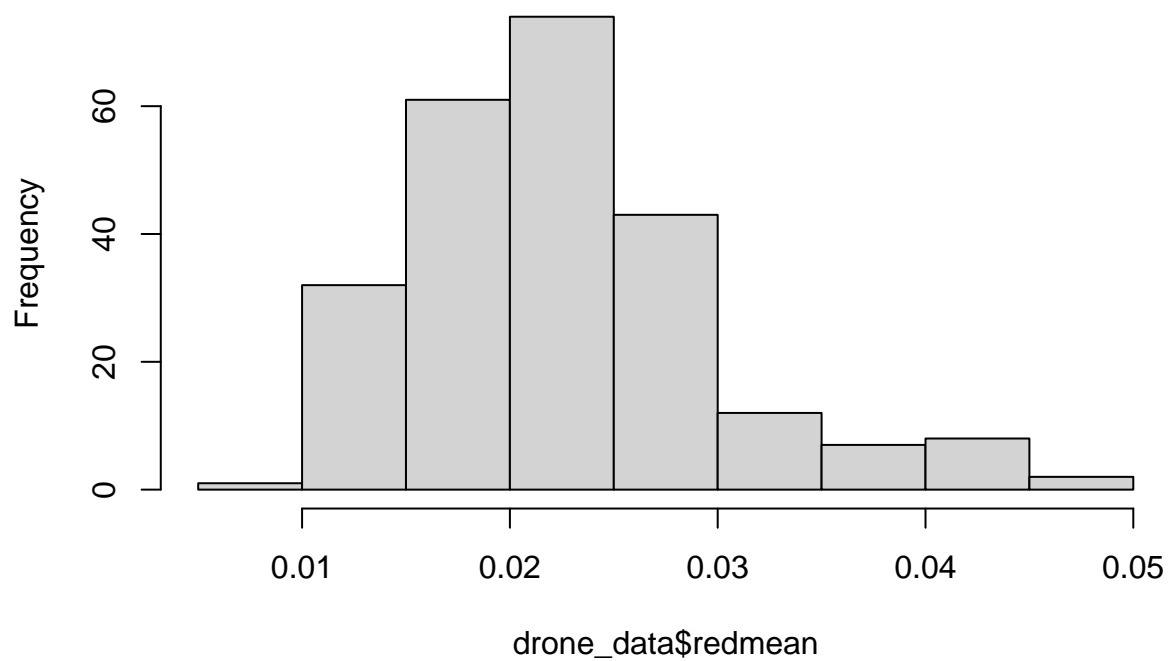


```
boxplot(drone_data$redmean)
```

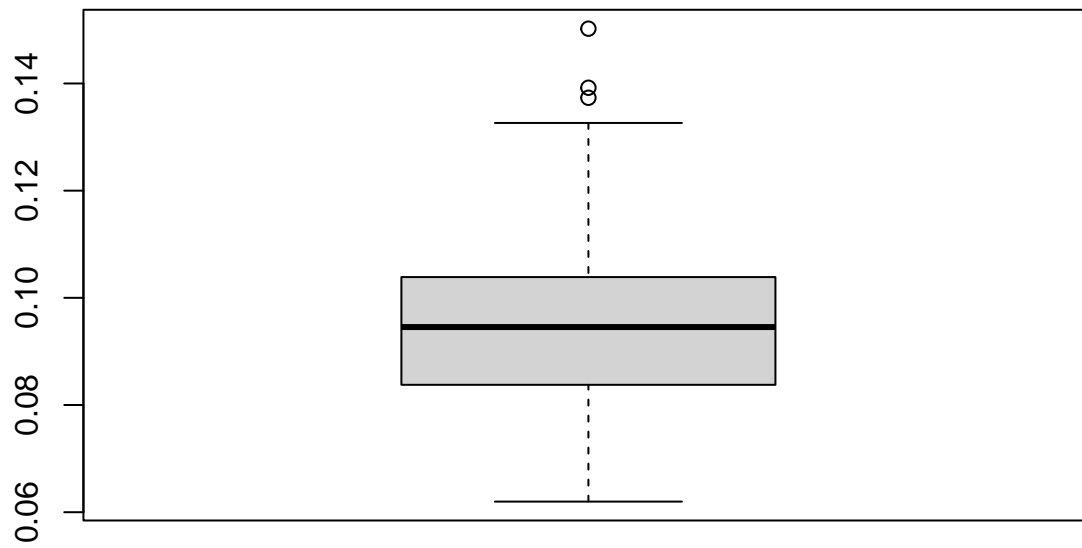



```
hist(drone_data$redmean)
```

Histogram of drone_data\$redmean

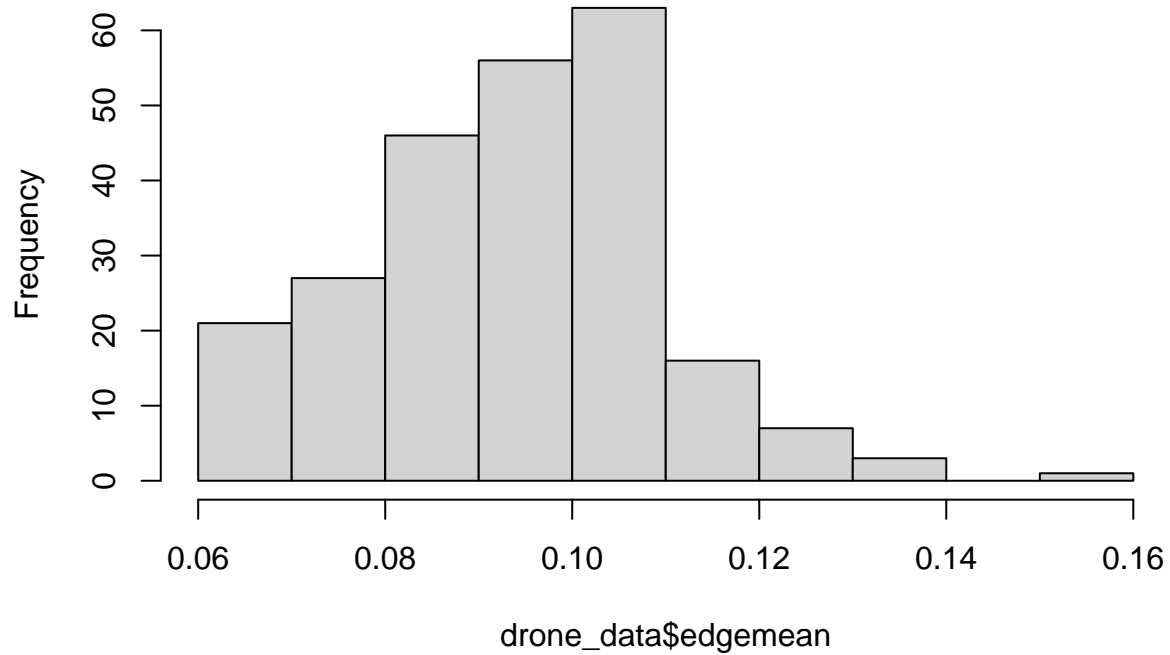


```
boxplot(drone_data$edgemean)
```

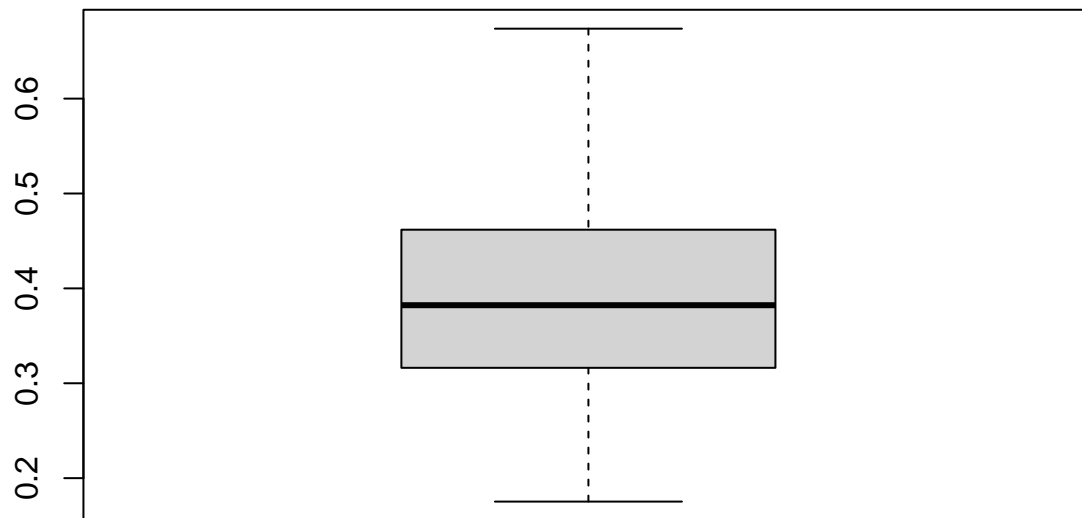


```
hist(drone_data$edgemean)
```

Histogram of drone_data\$edgemean

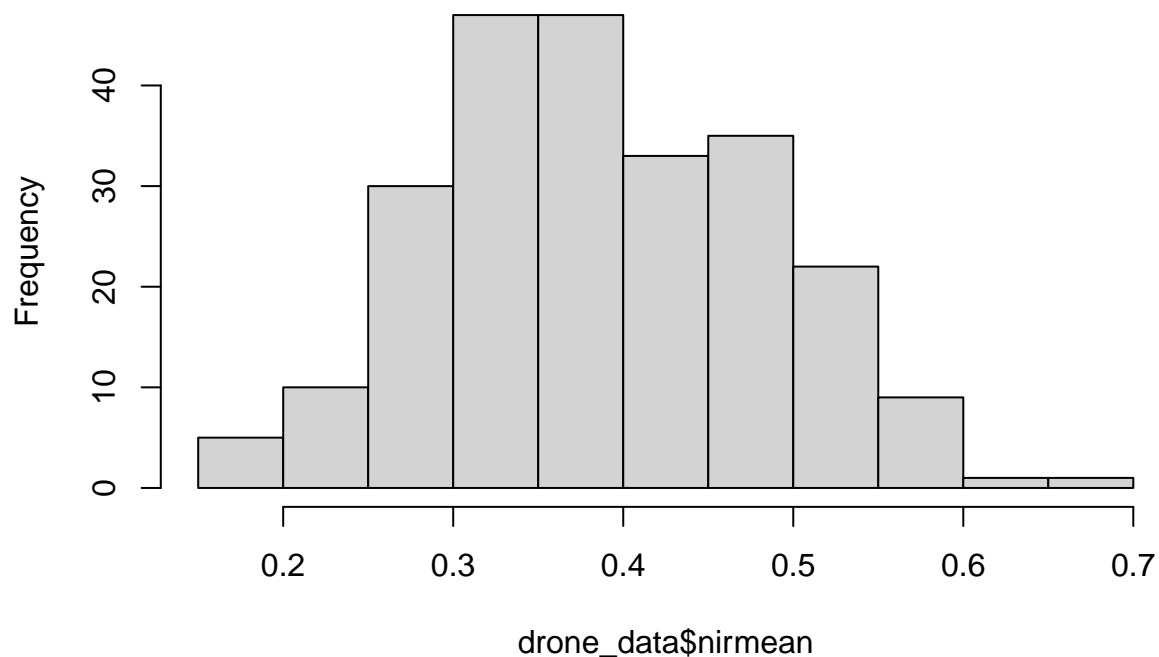


```
boxplot(drone_data$nirmean)
```



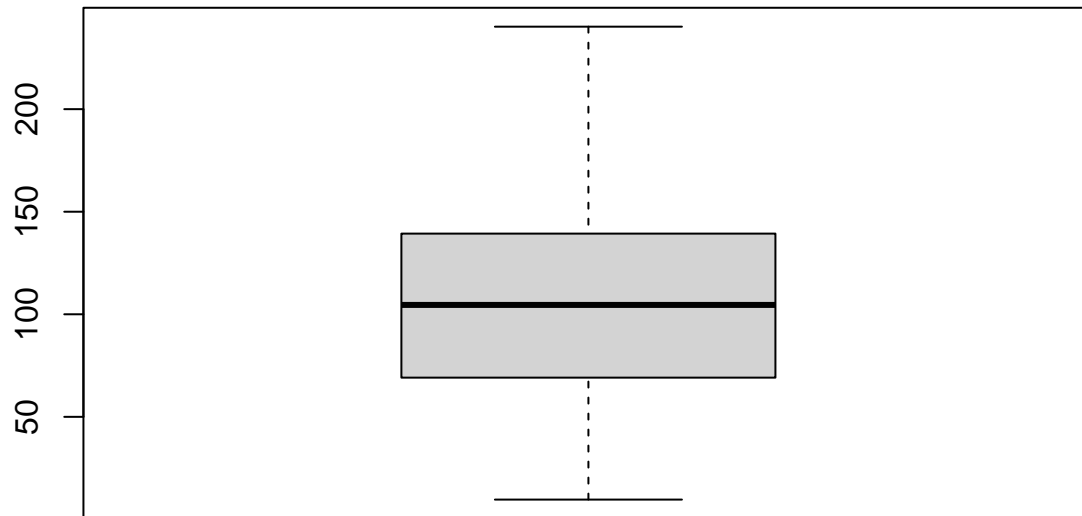
```
hist(drone_data$nirmean)
```

Histogram of drone_data\$nirmean



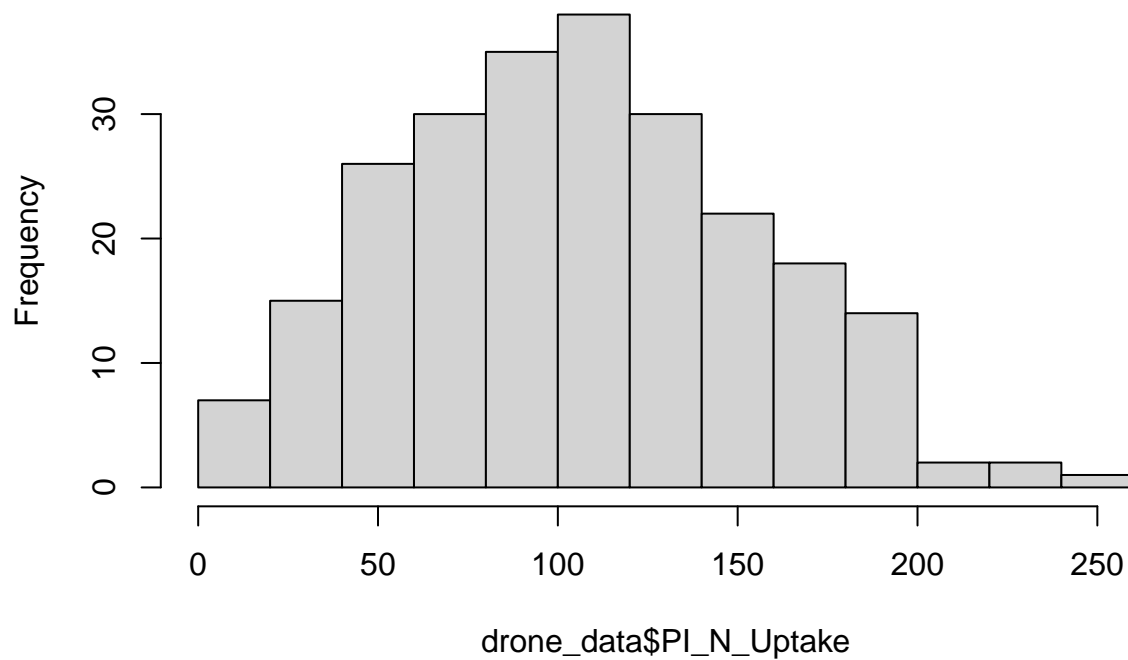
```
drone_data <- drone_data %>%
  mutate( biomass_dry_wt = biomass_plus_bag_g - (paper_bag_g * num_of_paper_bags) ,
          aboveground_biomass = (biomass_dry_wt / ring_size) * 10 , #ring size 0.5 m^2 biomass in kg per ha
          n_content = sample_N Ug / sample_weight_mg ,
          PI_N_Uptake = (aboveground_biomass * n_content) / 1000 #n uptake in kg per ha
          )#processes the data2

boxplot(drone_data$PI_N_Uptake)
```



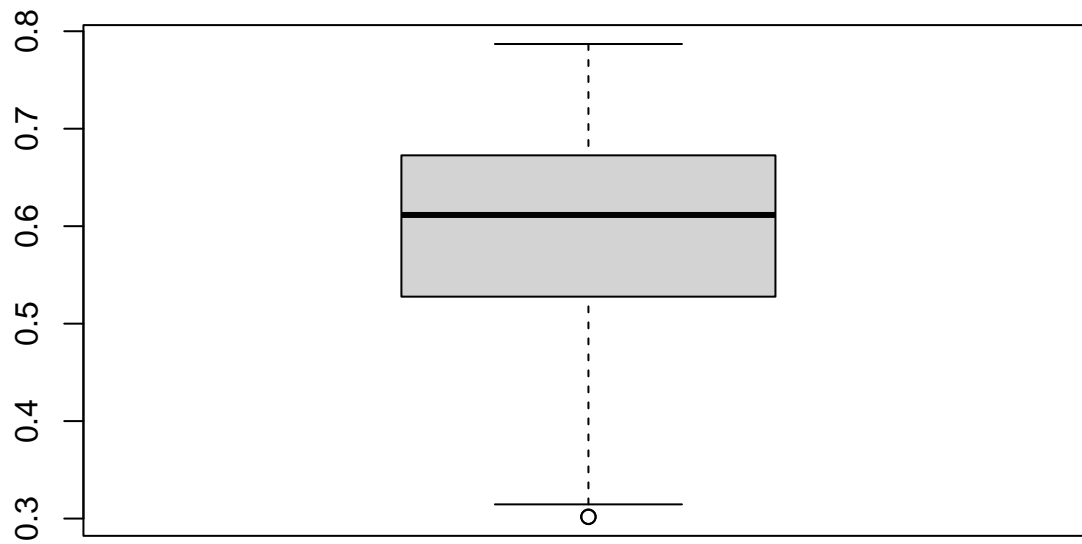
```
hist(drone_data$PI_N_Uptake)
```

Histogram of drone_data\$PI_N_Uptake



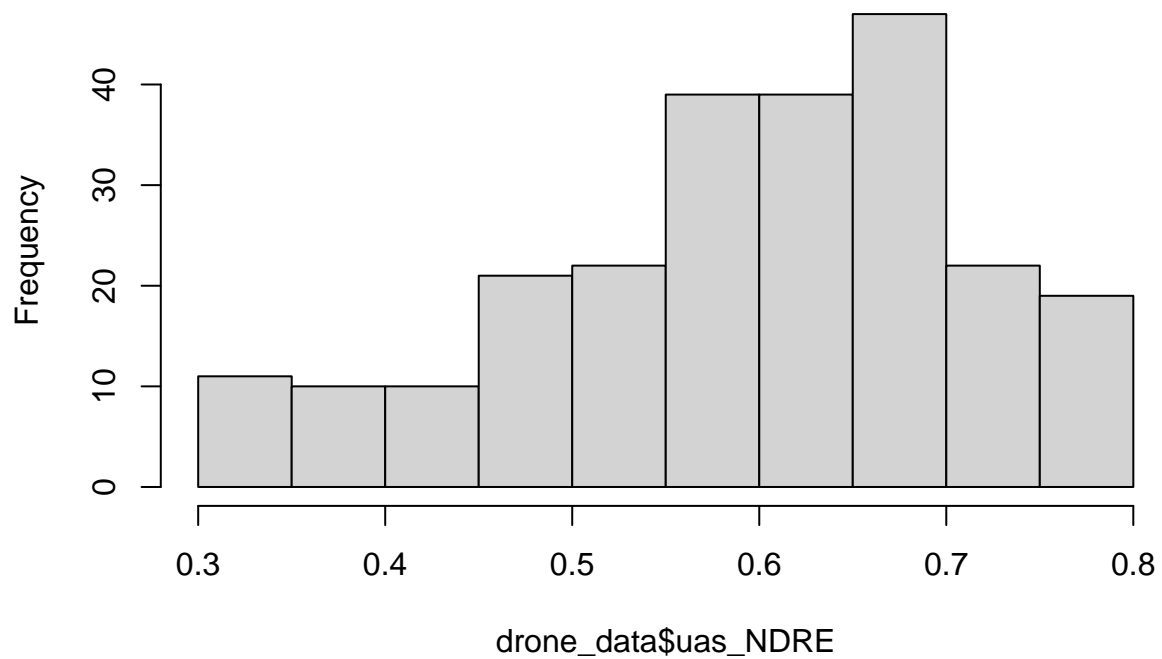
```
drone_data <- drone_data %>%
  mutate(uas_NDRE = ((nirmean - edgemean) / (nirmean + edgemean)) ,
         uas_NDVI = ((nirmean - redmean) / (nirmean + redmean))
         ) #calculates NDRE and NDVI

boxplot(drone_data$uas_NDRE)
```

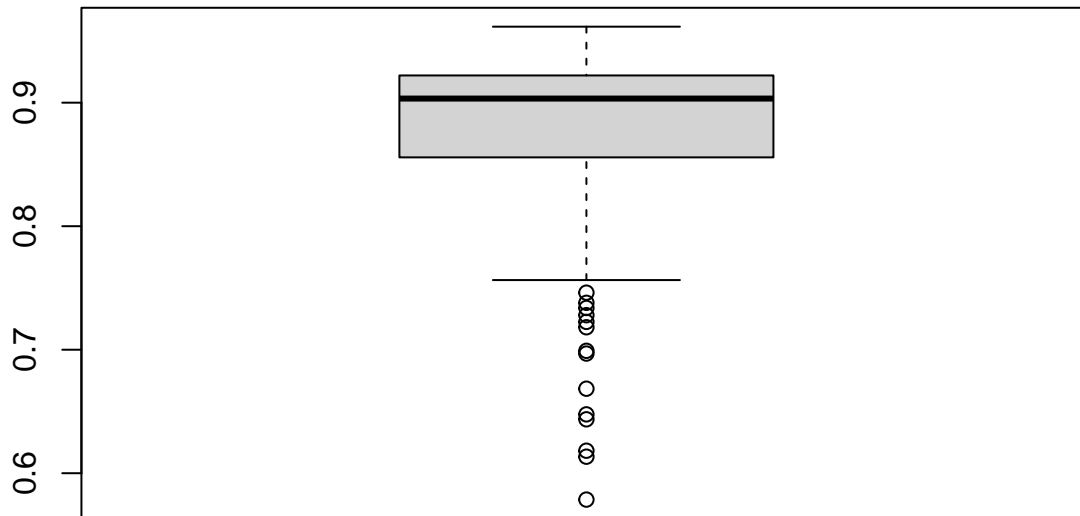


```
hist(drone_data$uas_NDRE)
```

Histogram of drone_data\$uas_NDRE

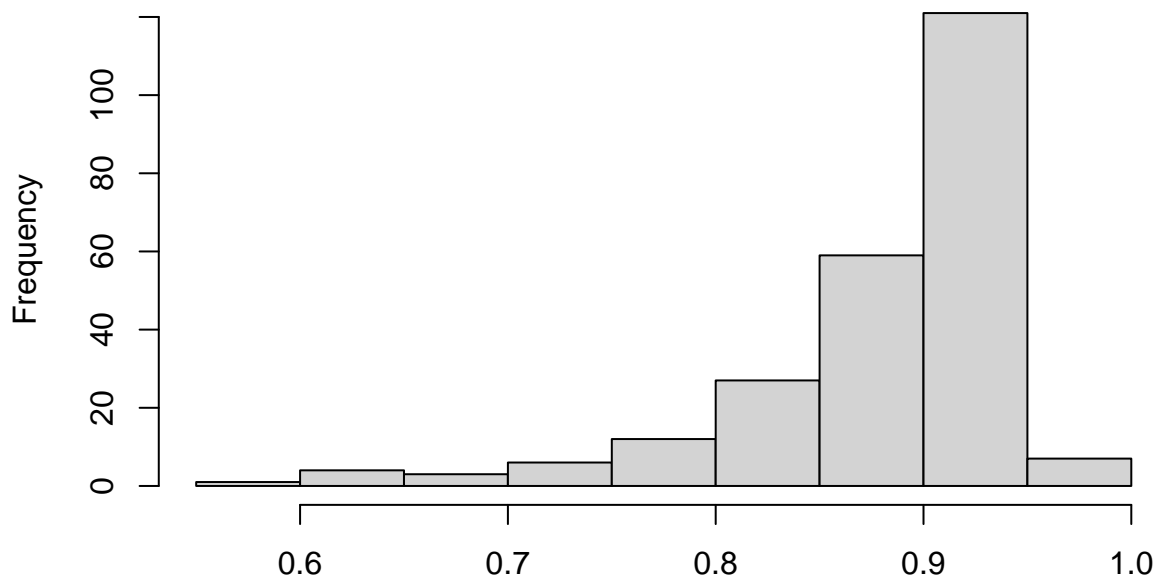


```
boxplot(drone_data$uas_NDVI)
```



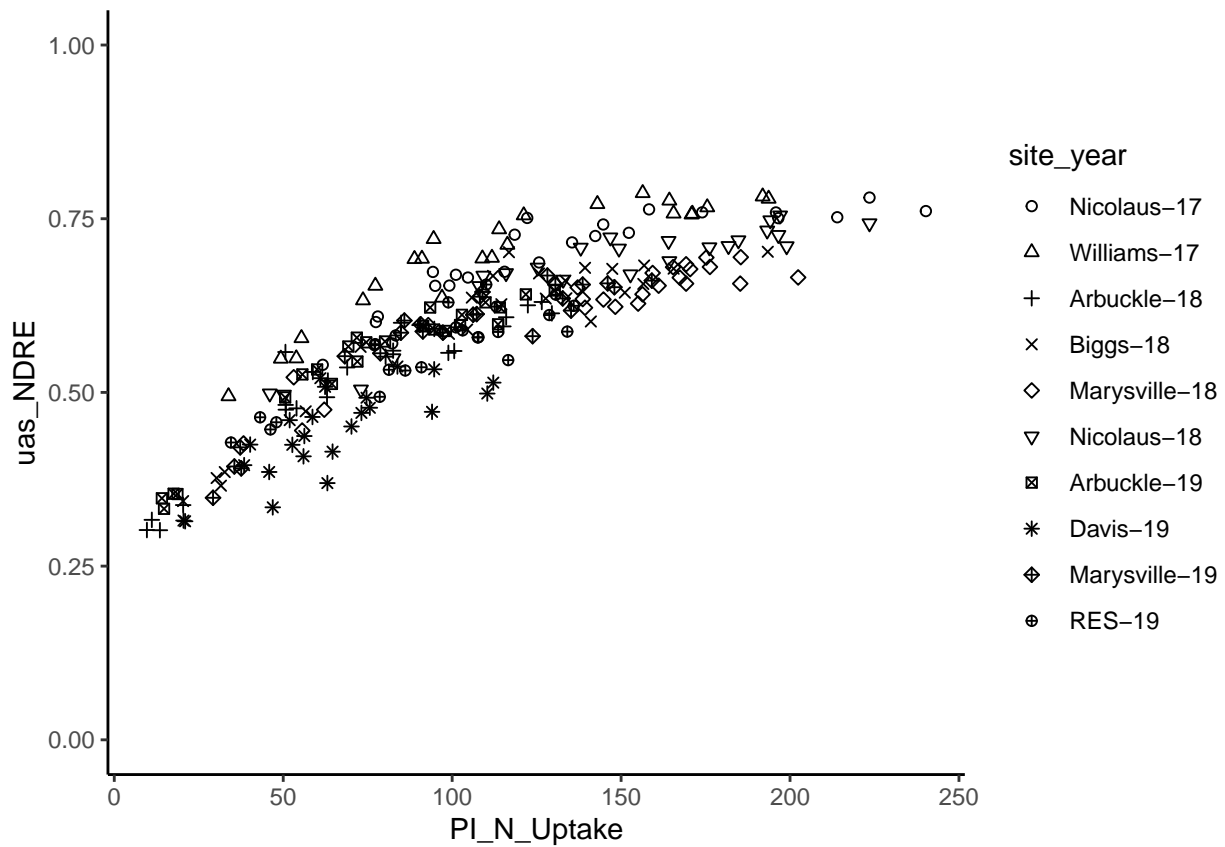
```
hist(drone_data$uas_NDVI)
```

Histogram of drone_data\$uas_NDVI

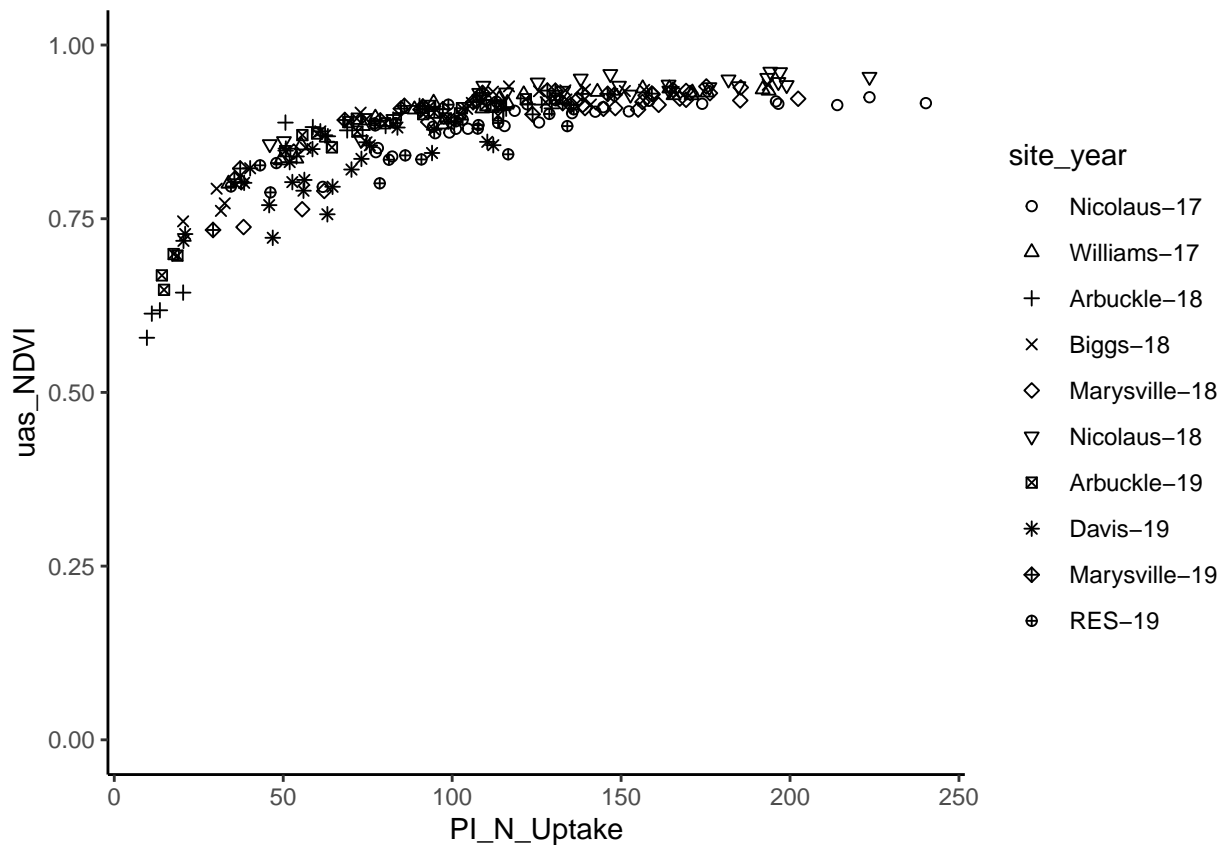


drone_data\$uas_NDVI

```
ggplot(data = drone_data, aes(x = PI_N_Uptake , y = uas_NDRE )) +
  geom_point(mapping = aes(x = PI_N_Uptake , y = uas_NDRE , shape = site_year) , data = drone_data) +
  theme_classic() +
  scale_shape_manual(values = seq(4:20)) +
  coord_cartesian(ylim = c(0,1))
```



```
ggplot(data = drone_data, aes(x = PI_N_Uptake , y = uas_NDVI )) +
  geom_point(mapping = aes(x = PI_N_Uptake , y = uas_NDVI , shape = site_year) , data = drone_data) +
  theme_classic() +
  scale_shape_manual(values = seq(4:20)) +
  coord_cartesian(ylim = c(0,1))
```



Calculating UAS RI

#gets the max NDRE value for each site

```
max_drone_data <- drone_data %>%
  filter(N_level_kgha_f %in% c(225, 235)) %>%
  dplyr::select(site_year, uas_NDVI, uas_NDRE) %>%
  group_by(site_year) %>%
  summarise_all(.funs = mean) %>%
  ungroup()

nic17 <- subset(max_drone_data, site_year == "Nicolaus-17")
nic17maxuas_NDRE <- as.numeric(nic17$uas_NDRE)
nic17maxuas_NDRE
```

```
## [1] 0.7642401
```

```
nic17maxuas_NDVI <- as.numeric(nic17$uas_NDVI)
nic17maxuas_NDVI
```

```
## [1] 0.9186193
```

```
wil17 <- subset(max_drone_data, site_year == "Williams-17")
wil17maxuas_NDRE <- as.numeric(wil17$uas_NDRE)
wil17maxuas_NDRE
```

```
## [1] 0.7785445
```



```

wil17maxuas_NDVI <- as.numeric(wil17$uas_NDVI)
wil17maxuas_NDVI

## [1] 0.9349728

arb18 <- subset(max_drone_data, site_year == "Arbuckle-18")
arb18maxuas_NDRE <- as.numeric(arb18$uas_NDRE)
arb18maxuas_NDRE

## [1] 0.6151996

arb18maxuas_NDVI <- as.numeric(arb18$uas_NDVI)
arb18maxuas_NDVI

## [1] 0.9113451

biggs18 <- subset(max_drone_data, site_year == "Biggs-18")
biggs18maxuas_NDRE <- as.numeric(biggs18$uas_NDRE)
biggs18maxuas_NDRE

## [1] 0.6796539

biggs18maxuas_NDVI <- as.numeric(biggs18$uas_NDVI)
biggs18maxuas_NDVI

## [1] 0.9342827

mry18 <- subset(max_drone_data, site_year == "Marysville-18")
mry18maxuas_NDRE <- as.numeric(mry18$uas_NDRE)
mry18maxuas_NDRE

## [1] 0.6812887

mry18maxuas_NDVI <- as.numeric(mry18$uas_NDVI)
mry18maxuas_NDVI

## [1] 0.9315267

nic18 <- subset(max_drone_data, site_year == "Nicolaus-18")
nic18maxuas_NDRE <- as.numeric(nic18$uas_NDRE)
nic18maxuas_NDRE

## [1] 0.7430412

nic18maxuas_NDVI <- as.numeric(nic18$uas_NDVI)
nic18maxuas_NDVI

## [1] 0.9558061

arb19 <- subset(max_drone_data, site_year == "Arbuckle-19")
arb19maxuas_NDRE <- as.numeric(arb19$uas_NDRE)
arb19maxuas_NDRE

## [1] 0.6288361

arb19maxuas_NDVI <- as.numeric(arb19$uas_NDVI)
arb19maxuas_NDVI

## [1] 0.9142775

```

```
davis19 <- subset(max_drone_data, site_year == "Davis-19")
davis19maxuas_NDRE <- as.numeric(davis19$uas_NDRE)
davis19maxuas_NDRE
```

```
## [1] 0.5012478
```

```
davis19maxuas_NDVI <- as.numeric(davis19$uas_NDVI)
davis19maxuas_NDVI
```

```
## [1] 0.8592071
```

```
mry19 <- subset(max_drone_data, site_year == "Marysville-19")
mry19maxuas_NDRE <- as.numeric(mry19$uas_NDRE)
mry19maxuas_NDRE
```

```
## [1] 0.655337
```

```
mry19maxuas_NDVI <- as.numeric(mry19$uas_NDVI)
mry19maxuas_NDVI
```

```
## [1] 0.9276236
```

```
res19 <- subset(max_drone_data, site_year == "RES-19")
res19maxuas_NDRE <- as.numeric(res19$uas_NDRE)
res19maxuas_NDRE
```

```
## [1] 0.5945887
```

```
res19maxuas_NDVI <- as.numeric(res19$uas_NDVI)
res19maxuas_NDVI
```

```
## [1] 0.8840389
```

```
drone_data <- drone_data %>%
```

```
  mutate(max_uas_NDRE = case_when(
    site_year == "Nicolaus-17" ~ nic17maxuas_NDRE ,
    site_year == "Williams-17" ~ wil17maxuas_NDRE ,
    site_year == "Arbuckle-18" ~ arb18maxuas_NDRE ,
    site_year == "Biggs-18" ~ biggs18maxuas_NDRE ,
    site_year == "Marysville-18" ~ mry18maxuas_NDRE ,
    site_year == "Nicolaus-18" ~ nic18maxuas_NDRE ,
    site_year == "Arbuckle-19" ~ arb19maxuas_NDRE ,
    site_year == "Davis-19" ~ davis19maxuas_NDRE ,
    site_year == "Marysville-19" ~ mry19maxuas_NDRE ,
    site_year == "RES-19" ~ res19maxuas_NDRE) #assign the max NDRE value for each
  )
```

```
drone_data <- drone_data %>%
```

```
  mutate(uas_NDRE_Response_Index = case_when(
    site_year == "Nicolaus-17" ~ nic17maxuas_NDRE / uas_NDRE,
    site_year == "Williams-17" ~ wil17maxuas_NDRE / uas_NDRE,
    site_year == "Arbuckle-18" ~ arb18maxuas_NDRE / uas_NDRE ,
    site_year == "Biggs-18" ~ biggs18maxuas_NDRE / uas_NDRE ,
    site_year == "Marysville-18" ~ mry18maxuas_NDRE / uas_NDRE ,
    site_year == "Nicolaus-18" ~ nic18maxuas_NDRE / uas_NDRE,
    site_year == "Arbuckle-19" ~ arb19maxuas_NDRE / uas_NDRE,
    site_year == "Davis-19" ~ davis19maxuas_NDRE / uas_NDRE,
    site_year == "Marysville-19" ~ mry19maxuas_NDRE / uas_NDRE,
  )
```

```

        site_year == "RES-19" ~ res19maxuas_NDRE / uas_NDRE
    )) #calculates uas_NDRE response index

drone_data <- drone_data %>%
  mutate(max_uas_NDVI = case_when(
    site_year == "Nicolaus-17" ~ nic17maxuas_NDVI ,
    site_year == "Williams-17" ~ will17maxuas_NDVI ,
    site_year == "Arbuckle-18" ~ arb18maxuas_NDVI ,
    site_year == "Biggs-18" ~ biggs18maxuas_NDVI ,
    site_year == "Marysville-18" ~ mry18maxuas_NDVI ,
    site_year == "Nicolaus-18" ~ nic18maxuas_NDVI ,
    site_year == "Arbuckle-19" ~ arb19maxuas_NDVI ,
    site_year == "Davis-19" ~ davis19maxuas_NDVI ,
    site_year == "Marysville-19" ~ mry19maxuas_NDVI ,
    site_year == "RES-19" ~ res19maxuas_NDVI) #assign max ndvi value for each sit
  )

drone_data <- drone_data %>%
  mutate(uas_NDVI_Response_Index = case_when(
    site_year == "Nicolaus-17" ~ nic17maxuas_NDVI / uas_NDVI,
    site_year == "Williams-17" ~ will17maxuas_NDVI / uas_NDVI,
    site_year == "Arbuckle-18" ~ arb18maxuas_NDVI / uas_NDVI ,
    site_year == "Biggs-18" ~ biggs18maxuas_NDVI / uas_NDVI ,
    site_year == "Marysville-18" ~ mry18maxuas_NDVI / uas_NDVI ,
    site_year == "Nicolaus-18" ~ nic18maxuas_NDVI / uas_NDVI,
    site_year == "Arbuckle-19" ~ arb19maxuas_NDVI / uas_NDVI,
    site_year == "Davis-19" ~ davis19maxuas_NDVI / uas_NDVI,
    site_year == "Marysville-19" ~ mry19maxuas_NDVI / uas_NDVI,
    site_year == "RES-19" ~ res19maxuas_NDVI / uas_NDVI
  )) #calculates uas_NDVI response index

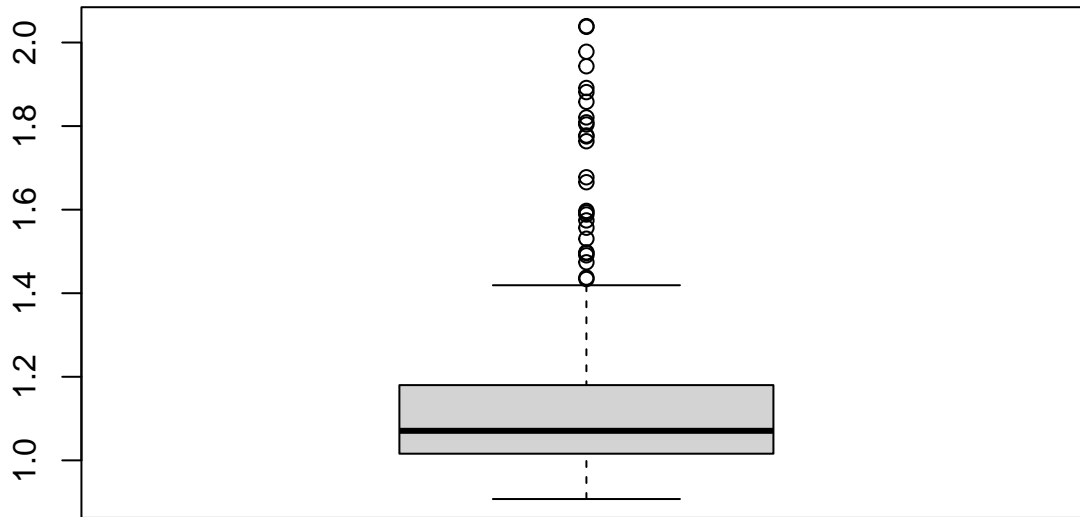
str(drone_data , give.attr = F)

## tibble [240 x 29] (S3: tbl_df/tbl/data.frame)
## $ site_year      : Factor w/ 10 levels "Nicolaus-17",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ year           : Factor w/ 3 levels "2017","2018",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ exp_plot_number : Factor w/ 27 levels "101","102","103",...: 1 2 3 4 6 7 9 10 11 12 ...
## $ Block          : Factor w/ 4 levels "1","2","3","4": 1 1 1 1 1 1 2 2 2 2 ...
## $ MainPlot       : Factor w/ 7 levels "1","2","3","4",...: 1 2 3 4 6 7 2 3 4 5 ...
## $ N_level        : Factor w/ 11 levels "0","45","75",...: 11 1 6 8 2 3 3 6 1 8 ...
## $ N_level_kgha   : num [1:240] 225 0 125 175 45 75 75 125 0 175 ...
## $ N_level_kgha_f : Factor w/ 11 levels "0","45","75",...: 10 1 5 8 2 3 3 5 1 8 ...
## $ biomass_plus_bag_g : num [1:240] 361 264 318 360 285 ...
## $ ring_size      : num [1:240] 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 ...
## $ paper_bag_g    : num [1:240] 45.5 45.5 45.5 45.5 45.5 45.5 45.5 45.5 45.5 45.5 ...
## $ num_of_paper_bags : num [1:240] 1 1 1 1 1 1 1 1 1 1 ...
## $ sample_weight_mg : num [1:240] 3.17 3.42 3.09 3.07 3.2 ...
## $ sample_N_ug     : num [1:240] 79.4 61.1 80.7 95.9 63.6 ...
## $ bluemean       : num [1:240] NA NA NA NA NA NA NA NA NA NA ...
## $ greenmean      : num [1:240] 0.0467 0.0581 0.0498 0.0488 0.0608 0.0589 0.0537 0.0488 0.05...
## $ redmean        : num [1:240] 0.023 0.0286 0.0243 0.0237 0.0292 0.0281 0.0263 0.0237 0.029...
## $ edgemean       : num [1:240] 0.0738 0.0867 0.0768 0.0764 0.0903 0.0877 0.0812 0.0754 0.08...
## $ nirmean        : num [1:240] 0.55 0.357 0.482 0.537 0.431 ...
## $ biomass_dry_wt  : num [1:240] 316 219 273 315 239 ...

```

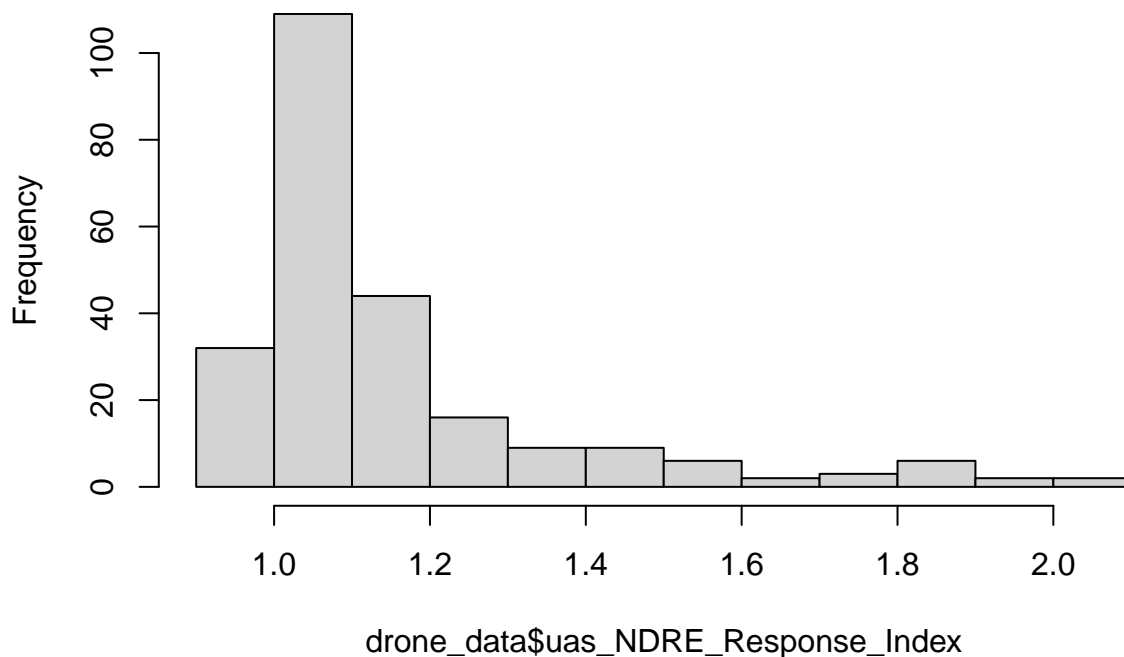
```
## $ aboveground_biomass : num [1:240] 6316 4376 5456 6296 4782 ...
## $ n_content           : num [1:240] 25.1 17.8 26.1 31.2 19.9 ...
## $ PI_N_Uptake         : num [1:240] 158 78 142 197 95 ...
## $ uas_NDRE            : num [1:240] 0.763 0.609 0.725 0.751 0.653 ...
## $ uas_NDVI            : num [1:240] 0.92 0.852 0.904 0.915 0.873 ...
## $ max_uas_NDRE        : num [1:240] 0.764 0.764 0.764 0.764 0.764 ...
## $ uas_NDRE_Response_Index: num [1:240] 1 1.25 1.05 1.02 1.17 ...
## $ max_uas_NDVI        : num [1:240] 0.919 0.919 0.919 0.919 0.919 ...
## $ uas_NDVI_Response_Index: num [1:240] 0.999 1.079 1.016 1.003 1.052 ...
```

```
boxplot(drone_data$uas_NDRE_Response_Index)
```

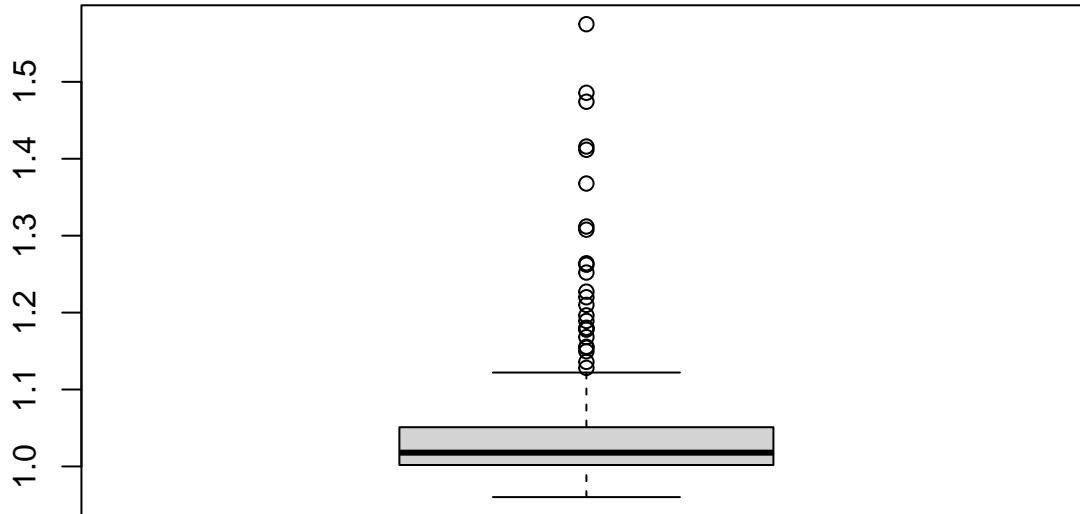


```
hist(drone_data$uas_NDRE_Response_Index)
```

Histogram of drone_data\$uas_NDRE_Response_Index

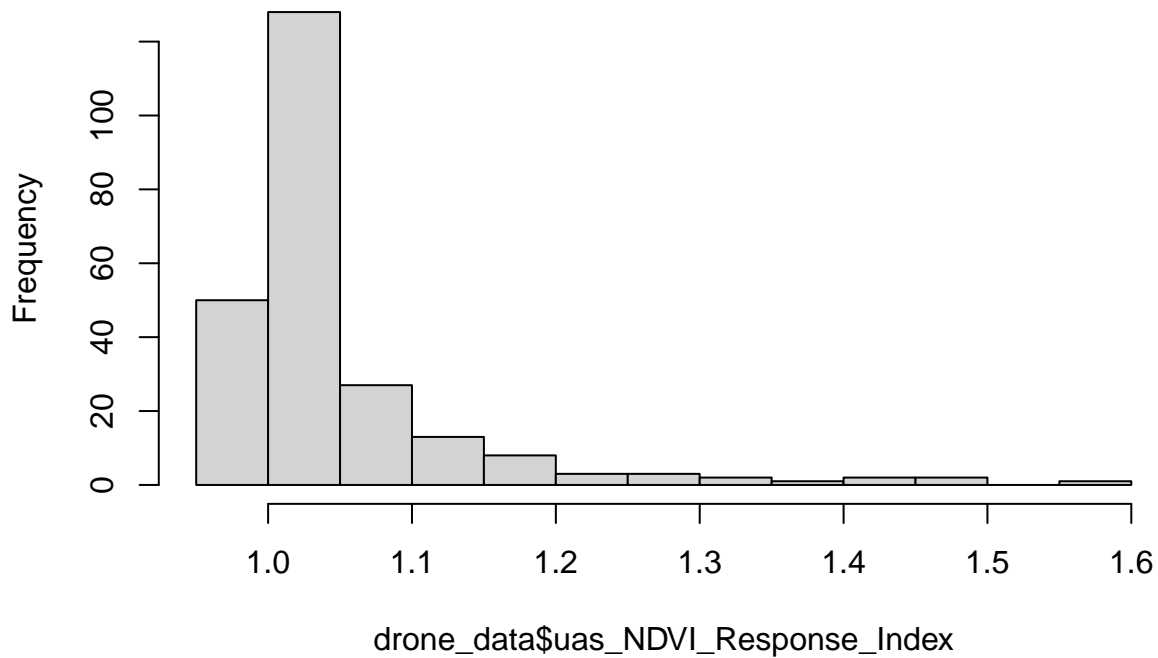


```
boxplot(drone_data$uas_NDVI_Response_Index)
```



```
hist(drone_data$uas_NDVI_Response_Index)
```

Histogram of drone_data\$uas_NDVI_Response_Index



```
drone_data <- dplyr::select(drone_data ,
  site_year,
  year,
  exp_plot_number,
  Block,
  MainPlot,
  N_level_kgha,
  N_level_kgha_f,
  PI_N_Uptake,
```

```

        uas_NDRE,
        uas_NDVI,
        uas_NDRE_Response_Index,
        uas_NDVI_Response_Index
    )#selects the relevant columns

sUAS_yield_data <- yield_data

sUAS_yield_data$site_year <- factor(sUAS_yield_data$site_year , levels = c("Nicolaus-17" , "Williams-17"

str(drone_data , give.attr = F)

## tibble [240 x 12] (S3: tbl_df/tbl/data.frame)
## $ site_year      : Factor w/ 10 levels "Nicolaus-17",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ year           : Factor w/ 3 levels "2017","2018",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ exp_plot_number : Factor w/ 27 levels "101","102","103",...: 1 2 3 4 6 7 9 10 11 12 ...
## $ Block          : Factor w/ 4 levels "1","2","3","4": 1 1 1 1 1 1 2 2 2 2 ...
## $ MainPlot        : Factor w/ 7 levels "1","2","3","4",...: 1 2 3 4 6 7 2 3 4 5 ...
## $ N_level_kgha     : num [1:240] 225 0 125 175 45 75 75 125 0 175 ...
## $ N_level_kgha_f   : Factor w/ 11 levels "0","45","75",...: 10 1 5 8 2 3 3 5 1 8 ...
## $ PI_N_Uptake      : num [1:240] 158 78 142 197 95 ...
## $ uas_NDRE         : num [1:240] 0.763 0.609 0.725 0.751 0.653 ...
## $ uas_NDVI         : num [1:240] 0.92 0.852 0.904 0.915 0.873 ...
## $ uas_NDRE_Response_Index: num [1:240] 1 1.25 1.05 1.02 1.17 ...
## $ uas_NDVI_Response_Index: num [1:240] 0.999 1.079 1.016 1.003 1.052 ...

str(sUAS_yield_data , give.attr = F)

## tibble [240 x 11] (S3: tbl_df/tbl/data.frame)
## $ site_year      : Factor w/ 10 levels "Nicolaus-17",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ year           : Factor w/ 3 levels "2017","2018",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ Block          : Factor w/ 4 levels "1","2","3","4": 1 1 1 1 1 1 2 2 2 2 ...
## $ MainPlot        : Factor w/ 7 levels "1","2","3","4",...: 1 2 3 4 6 7 2 3 4 5 ...
## $ SubPlot         : Factor w/ 3 levels "1","2","3": 2 3 1 1 1 2 1 1 2 3 ...
## $ exp_plot_number: Factor w/ 27 levels "101","102","103",...: 1 2 3 4 6 7 9 10 11 12 ...
## $ N_level_kgha     : num [1:240] 225 0 125 175 45 75 75 125 0 175 ...
## $ N_level_kgha_f   : Factor w/ 11 levels "0","45","75",...: 10 1 5 8 2 3 3 5 1 8 ...
## $ TopDress_kgha    : num [1:240] 0 0 0 0 0 0 0 0 0 0 ...
## $ TopDress_kgha_f  : Factor w/ 1 level "0": 1 1 1 1 1 1 1 1 1 1 ...
## $ GrainYield_Mgha : num [1:240] 11.4 11 12 11.9 11.5 ...

sUAS_data <- full_join( drone_data , sUAS_yield_data)

## Joining, by = c("site_year", "year", "exp_plot_number", "Block", "MainPlot",
## "N_level_kgha", "N_level_kgha_f")

sUAS_data <- dplyr::select(sUAS_data ,
    site_year,
    year,
    exp_plot_number,
    Block,
    MainPlot,
    SubPlot,
    N_level_kgha,
    N_level_kgha_f,
    TopDress_kgha,

```

```

TopDress_kgha_f,
PI_N_Uptake,
uas_NDRE,
uas_NDVI,
GrainYield_Mgha,
uas_NDRE_Response_Index,
uas_NDVI_Response_Index
)#selects the relevant columns

str(sUAS_data , give.attr = F)

## tibble [240 x 16] (S3: tbl_df/tbl/data.frame)
## $ site_year      : Factor w/ 10 levels "Nicolaus-17",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ year           : Factor w/ 3 levels "2017","2018",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ exp_plot_number : Factor w/ 27 levels "101","102","103",...: 1 2 3 4 6 7 9 10 11 12 ...
## $ Block          : Factor w/ 4 levels "1","2","3","4": 1 1 1 1 1 1 2 2 2 2 ...
## $ MainPlot        : Factor w/ 7 levels "1","2","3","4",...: 1 2 3 4 6 7 2 3 4 5 ...
## $ SubPlot         : Factor w/ 3 levels "1","2","3": 2 3 1 1 1 2 1 1 2 3 ...
## $ N_level_kgha    : num [1:240] 225 0 125 175 45 75 75 125 0 175 ...
## $ N_level_kgha_f  : Factor w/ 11 levels "0","45","75",...: 10 1 5 8 2 3 3 5 1 8 ...
## $ TopDress_kgha   : num [1:240] 0 0 0 0 0 0 0 0 0 0 ...
## $ TopDress_kgha_f : Factor w/ 1 level "0": 1 1 1 1 1 1 1 1 1 1 ...
## $ PI_N_Uptake     : num [1:240] 158 78 142 197 95 ...
## $ uas_NDRE        : num [1:240] 0.763 0.609 0.725 0.751 0.653 ...
## $ uas_NDVI        : num [1:240] 0.92 0.852 0.904 0.915 0.873 ...
## $ GrainYield_Mgha : num [1:240] 11.4 11 12 11.9 11.5 ...
## $ uas_NDRE_Response_Index: num [1:240] 1 1.25 1.05 1.02 1.17 ...
## $ uas_NDVI_Response_Index: num [1:240] 0.999 1.079 1.016 1.003 1.052 ...

```

Combining the data

```

gs_data <- gs_data %>%
  mutate(gs_NDVI_SI = 1 / gs_NDVI_Response_Index)

sUAS_data <- sUAS_data %>%
  mutate(uas_NDVI_SI = 1 / uas_NDVI_Response_Index)

sUAS_data <- sUAS_data %>%
  mutate(uas_NDRE_SI = 1 / uas_NDRE_Response_Index)

gs_data <- gs_data %>%
  dplyr::select(site_year ,
                year,
                exp_plot_number,
                Block,
                MainPlot,
                SubPlot,
                N_level_kgha,
                N_level_kgha_f,
                TopDress_kgha,
                TopDress_kgha_f,
                PI_N_Uptake,

```

```

        NDVI,
        GrainYield_Mgha,
        gs_NDVI_Response_Index,
        gs_NDVI_SI
    )

sUAS_data <- sUAS_data %>%
  dplyr::select(site_year ,
    year,
    exp_plot_number,
    Block,
    MainPlot,
    SubPlot,
    N_level_kgha,
    N_level_kgha_f,
    TopDress_kgha,
    TopDress_kgha_f,
    uas_NDRE,
    uas_NDVI,
    uas_NDVI_Response_Index,
    uas_NDVI_SI,
    uas_NDRE_Response_Index,
    uas_NDRE_SI
  )

paper3_data <- full_join(gs_data , sUAS_data)

## Joining, by = c("site_year", "year", "exp_plot_number", "Block", "MainPlot",
## "SubPlot", "N_level_kgha", "N_level_kgha_f", "TopDress_kgha",
## "TopDress_kgha_f")

paper3_gsdata <- paper3_data %>%
  dplyr::select(site_year ,
    year ,
    exp_plot_number ,
    Block ,
    MainPlot ,
    SubPlot ,
    N_level_kgha ,
    N_level_kgha_f ,
    TopDress_kgha ,
    TopDress_kgha_f ,
    NDVI,
    gs_NDVI_Response_Index,
    gs_NDVI_SI ,
    GrainYield_Mgha,
    PI_N_Uptake) %>%
  mutate(Platform = "GreenSeeker NDVI" ) %>%
  rename(Index = NDVI ,
    SI = gs_NDVI_SI,
    RI = gs_NDVI_Response_Index)

paper3_uas_ndvi_data <- paper3_data %>%
  dplyr::select(site_year ,

```



```

        year ,
        exp_plot_number ,
        Block ,
        MainPlot ,
        SubPlot ,
        N_level_kgha ,
        N_level_kgha_f ,
        TopDress_kgha ,
        TopDress_kgha_f ,
        uas_NDVI,
        uas_NDVI_Response_Index,
        uas_NDVI_SI ,
        GrainYield_Mgha,
        PI_N_Uptake) %>%
mutate(Platform = "sUAS NDVI" ) %>%
rename(Index = uas_NDVI,
        SI = uas_NDVI_SI,
        RI = uas_NDVI_Response_Index)

paper3_uas_ndre_data <- paper3_data %>%
  dplyr::select(site_year ,
                year ,
                exp_plot_number ,
                Block ,
                MainPlot ,
                SubPlot ,
                N_level_kgha ,
                N_level_kgha_f ,
                TopDress_kgha ,
                TopDress_kgha_f ,
                uas_NDRE ,
                uas_NDRE_Response_Index ,
                uas_NDRE_SI ,
                GrainYield_Mgha,
                PI_N_Uptake) %>%
mutate(Platform = "sUAS NDRE" ) %>%
rename(Index = uas_NDRE ,
        SI = uas_NDRE_SI,
        RI = uas_NDRE_Response_Index)

paper3_data <- rbind(paper3_gsdata ,
                     paper3_uas_ndvi_data ,
                     paper3_uas_ndre_data)

paper3_data <- paper3_data %>%
  dplyr::select(site_year ,
                year ,
                Platform,
                exp_plot_number ,
                Block ,
                MainPlot ,
                SubPlot ,
                N_level_kgha ,
                N_level_kgha_f ,

```

```

TopDress_kgha ,
TopDress_kgha_f ,
Index,
RI,
SI ,
GrainYield_Mgha,
PI_N_Uptake)

paper3_data$Platform <- as.factor(paper3_data$Platform)

str(paper3_data , give.attr = F)

## rowwise_df [720 x 16] (S3: rowwise_df/tbl_df/tbl/data.frame)
## $ site_year      : Factor w/ 10 levels "Nicolaus-17",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ year           : Factor w/ 3 levels "2017","2018",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ Platform       : Factor w/ 3 levels "GreenSeeker NDVI",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ exp_plot_number: Factor w/ 27 levels "101","102","103",...: 1 2 3 4 6 7 9 10 11 12 ...
## $ Block          : Factor w/ 4 levels "1","2","3","4": 1 1 1 1 1 1 2 2 2 2 ...
## $ MainPlot       : Factor w/ 7 levels "1","2","3","4",...: 1 2 3 4 6 7 2 3 4 5 ...
## $ SubPlot        : Factor w/ 3 levels "1","2","3": 2 3 1 1 1 2 1 1 2 3 ...
## $ N_level_kgha    : num [1:720] 225 0 125 175 45 75 75 125 0 175 ...
## $ N_level_kgha_f  : Factor w/ 11 levels "0","45","75",...: 10 1 5 8 2 3 3 5 1 8 ...
## $ TopDress_kgha   : num [1:720] 0 0 0 0 0 0 0 0 0 0 ...
## $ TopDress_kgha_f: Factor w/ 1 level "0": 1 1 1 1 1 1 1 1 1 1 ...
## $ Index           : num [1:720] 0.777 0.493 0.683 0.787 0.683 ...
## $ RI              : num [1:720] 1.012 1.593 1.15 0.999 1.15 ...
## $ SI              : num [1:720] 0.988 0.628 0.87 1.001 0.87 ...
## $ GrainYield_Mgha: num [1:720] 11.4 11 12 11.9 11.5 ...
## $ PI_N_Uptake     : num [1:720] 158 78 142 197 95 ...

paper3_data <- tibble::rowid_to_column(paper3_data, "ID") #adds a columns with row number.

```

Calculating Relative Yield

```

nic17 <- subset(paper3_data, site_year == "Nicolaus-17")
nic17 <- apply(nic17,2,max)
nic17max_yield <- as.numeric(nic17[16])

wil17 <- subset(paper3_data, site_year == "Williams-17")
wil17 <- apply(wil17,2,max)
wil17max_yield <- as.numeric(wil17[16])

arb18 <- subset(paper3_data, site_year == "Arbuckle-18")
arb18 <- apply(arb18,2,max)
arb18max_yield <- as.numeric(arb18[16])

nic18 <- subset(paper3_data, site_year == "Nicolaus-18")
nic18 <- apply(nic18,2,max)
nic18max_yield <- as.numeric(nic18[16])

mry18 <- subset(paper3_data, site_year == "Marysville-18")
mry18 <- apply(mry18,2,max)
mry18max_yield <- as.numeric(mry18[16])

```

```

biggs18 <- subset(paper3_data, site_year == "Biggs-18")
biggs18 <- apply(biggs18,2,max)
biggs18max_yield <- as.numeric(biggs18[16])

arb19 <- subset(paper3_data, site_year == "Arbuckle-19")
arb19 <- apply(arb19,2,max)
arb19max_yield <- as.numeric(arb19[16])

dav19 <- subset(paper3_data, site_year == "Davis-19")
dav19 <- apply(dav19,2,max)
dav19max_yield <- as.numeric(dav19[16])

mry19 <- subset(paper3_data, site_year == "Marysville-19")
mry19 <- apply(mry19,2,max)
mry19max_yield <- as.numeric(mry19[16])

res19 <- subset(paper3_data, site_year == "RES-19")
res19 <- apply(res19,2,max)
res19max_yield <- as.numeric(res19[16])

paper3_data <- paper3_data %>%
  mutate(relative_grain_yield = case_when(
    site_year == "Nicolaus-17" ~ GrainYield_Mgha / nic17max_yield ,
    site_year == "Williams-17" ~ GrainYield_Mgha / will17max_yield ,
    site_year == "Arbuckle-18" ~ GrainYield_Mgha / arb18max_yield ,
    site_year == "Nicolaus-18" ~ GrainYield_Mgha / nic18max_yield ,
    site_year == "Marysville-18" ~ GrainYield_Mgha / mry18max_yield ,
    site_year == "Biggs-18" ~ GrainYield_Mgha / biggs18max_yield ,
    site_year == "Arbuckle-19" ~ GrainYield_Mgha / arb19max_yield ,
    site_year == "Davis-19" ~ GrainYield_Mgha / dav19max_yield ,
    site_year == "Marysville-19" ~ GrainYield_Mgha / mry19max_yield ,
    site_year == "RES-19" ~ GrainYield_Mgha / res19max_yield))

```

Outlier Removal

```

filter1 <- paper3_data %>%
  filter(site_year == "Biggs-18" & exp_plot_number == 101)

paper3_data <- paper3_data %>%
  filter(!ID %in% filter1$ID) #removes Biggs-18 plot 101 plot bc tractor ran through it

paper3_data$Platform = factor(paper3_data$Platform, levels=c( "GreenSeeker NDVI" , "sUAS NDRE" , "sUAS

paper3_data$RI[paper3_data$RI < 1] <- 1 #converts values less than 1, to 1.
paper3_data$SI[paper3_data$SI > 1] <- 1 #converts values less than 1, to 1.

hist_data <- paper3_data %>%
  dplyr::select(Platform , RI) %>%
  group_by(Platform) %>%
  summarise(mean = mean(RI) , sd = sd(RI) , median = median(RI))

hist_data_gs <- hist_data %>%

```

```

  filter(Platform == "GreenSeeker NDVI")

upper_limit <- hist_data_gs[2] + (5*hist_data_gs[3]) #upper limit for outlier removal. Observations tha

print(upper_limit)

##          mean
## 1 3.690994
#need to remove Arbuckle-18 plot 203 and Arbuckle-19 plot 104

filter1 <- paper3_data %>%
  filter(site_year == "Arbuckle-18" & exp_plot_number == 203)

filter2 <- paper3_data %>%
  filter(site_year == "Arbuckle-19" & exp_plot_number == 104)

filter <- rbind(filter1 , filter2)

filter$ID

## [1]  57 297 537 148 388 628

paper3_data <- paper3_data %>%
  filter(!ID %in% filter$ID)

paper3_gsdata <- paper3_data %>%
  filter(Platform == "GreenSeeker NDVI")

paper3_uas_ndvi_data <- paper3_data %>%
  filter(Platform == "sUAS NDVI")

paper3_uas_ndre_data <- paper3_data %>%
  filter(Platform == "sUAS NDRE")

```

FIGURE 2

Data

```

suppl_data <- paper3_data

suppl_data1 <- suppl_data %>%
  dplyr::select(site_year, N_level_kgha , TopDress_kgha_f , GrainYield_Mgha , PI_N_Uptake) %>%
  group_by(site_year , N_level_kgha ,TopDress_kgha_f ) %>%
  summarise_all(mean) %>%
  mutate(PI_N_Uptake_r = round(PI_N_Uptake , digits = 0),
         GrainYield_Mgha_r = round(GrainYield_Mgha , digits = 1)) %>%
  ungroup()

factor_list <- list( site = suppl_data1$site_year)

tapply(suppl_data1$GrainYield_Mgha_r, factor_list , max)

## site
##  Nicolaus-17  Williams-17  Arbuckle-18  Biggs-18 Marysville-18

```

```
##      12.2      11.1      12.7      12.3      11.3
## Nicolaus-18 Arbuckle-19 Davis-19 Marysville-19 RES-19
##      13.3      11.5      10.7      10.7      9.1
```

```
tapply(suppl_data1$GrainYield_Mgha_r, factor_list , min)
```

```
## site
## Nicolaus-17 Williams-17 Arbuckle-18 Biggs-18 Marysville-18
##      10.6      6.6      3.2      7.3      8.7
## Nicolaus-18 Arbuckle-19 Davis-19 Marysville-19 RES-19
##      9.2      4.2      7.6      6.9      6.0
```

Legend

```
suppl_data1_nic17 <- suppl_data1 %>%
  filter(site_year == "Nicolaus-17")
```

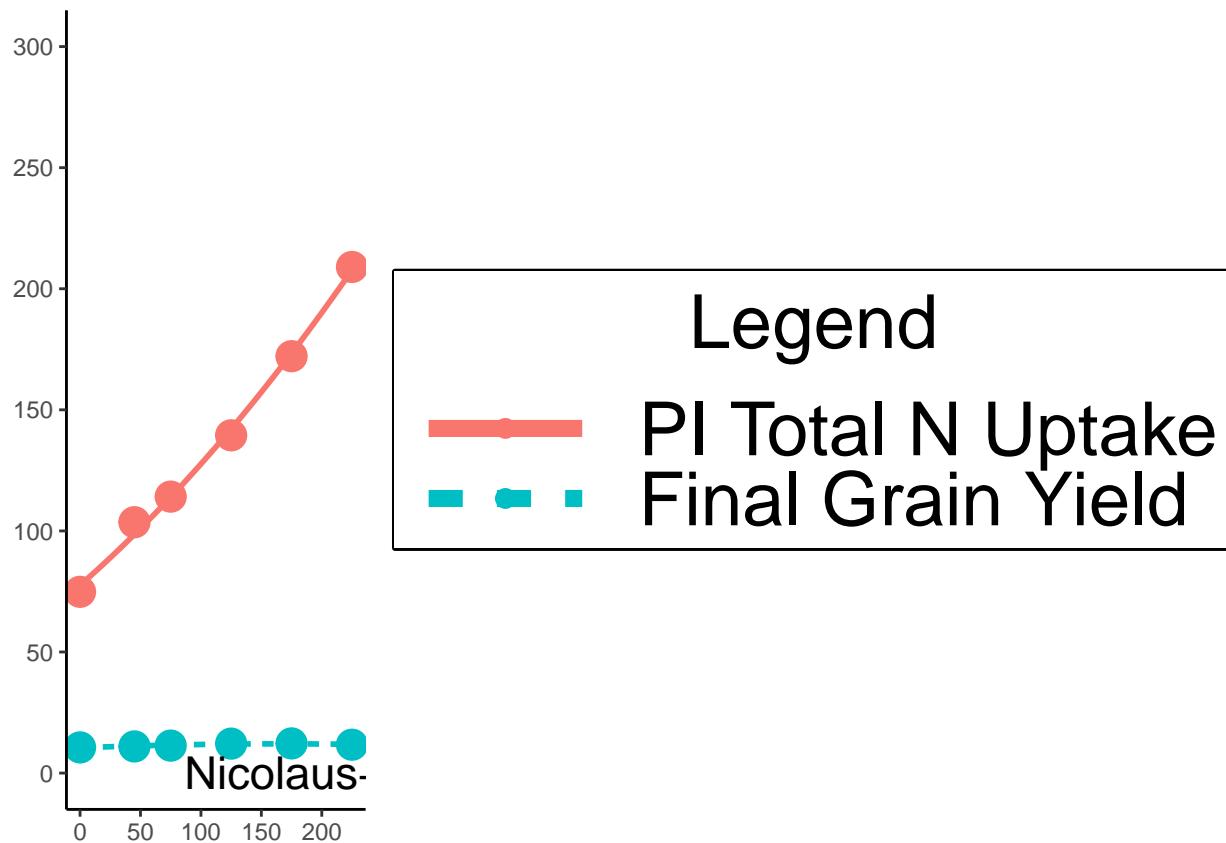
```
suppl_data1_nic17_yield <- suppl_data1_nic17 %>%
  dplyr::select(-c(PI_N_Uptake)) %>%
  mutate(Legend = as.factor("Final Grain Yield")) %>%
  rename("Unit" = GrainYield_Mgha)
```

```
suppl_data1_nic17_nup <- suppl_data1_nic17 %>%
  dplyr::select(-c(GrainYield_Mgha)) %>%
  mutate(Legend = as.factor("PI Total N Uptake")) %>%
  rename("Unit" = PI_N_Uptake)
```

```
suppl_data1_nic17 <- rbind(suppl_data1_nic17_nup , suppl_data1_nic17_yield)
```

```
nic17_plot_legend <- ggplot(data = suppl_data1_nic17 , aes( x = N_level_kgha , y = Unit , color = Legend)) +
  geom_point(data = suppl_data1_nic17 , aes ( x = N_level_kgha , y = Unit , color = Legend) , size = 5) +
  scale_y_continuous(breaks = seq(0 , 300 , by = 50)) +
  coord_cartesian(ylim = c(0 , 300)) +
  theme_classic() +
  theme(legend.text = element_text(size = 28 ),
        legend.title = element_text(size = 28 ),
        legend.box.background = element_rect(size = 1)) +
  annotate("text" , x = 190 , y = 0 , label = "Nicolaus-17" , size = 6) +
  labs( x = NULL , y = NULL) +
  geom_smooth(method = "lm", formula = y ~ poly(x, 2), se = FALSE) +
  guides(size = "none" , color = guide_legend(keywidth = 5 , keyheight = 1.5 , unit = "cm" , override.aes = FALSE))
```

```
nic17_plot_legend
```



```
legend <- get_legend(nic17_plot_legend)
```

Nicolaus-17

```
suppl_data1_nic17 <- suppl_data1 %>%
  filter(site_year == "Nicolaus-17")

nic17_nuptake_lm <- lm(PI_N_Uptake ~ N_level_kgha + I(N_level_kgha*N_level_kgha) , data = suppl_data1_nic17)

nic17_fitted <- fitted(nic17_nuptake_lm)
nic17_df <- data.frame(suppl_data1_nic17$N_level_kgha , nic17_fitted) #creates dataframe

nic17_yield_lm <- lm(GrainYield_Mgha ~ N_level_kgha + I(N_level_kgha*N_level_kgha) , data = suppl_data1_nic17)

nic17_fitted_2 <- (fitted(nic17_yield_lm)*23)
nic17_df_2 <- data.frame(suppl_data1_nic17$N_level_kgha , nic17_fitted_2) #creates dataframe

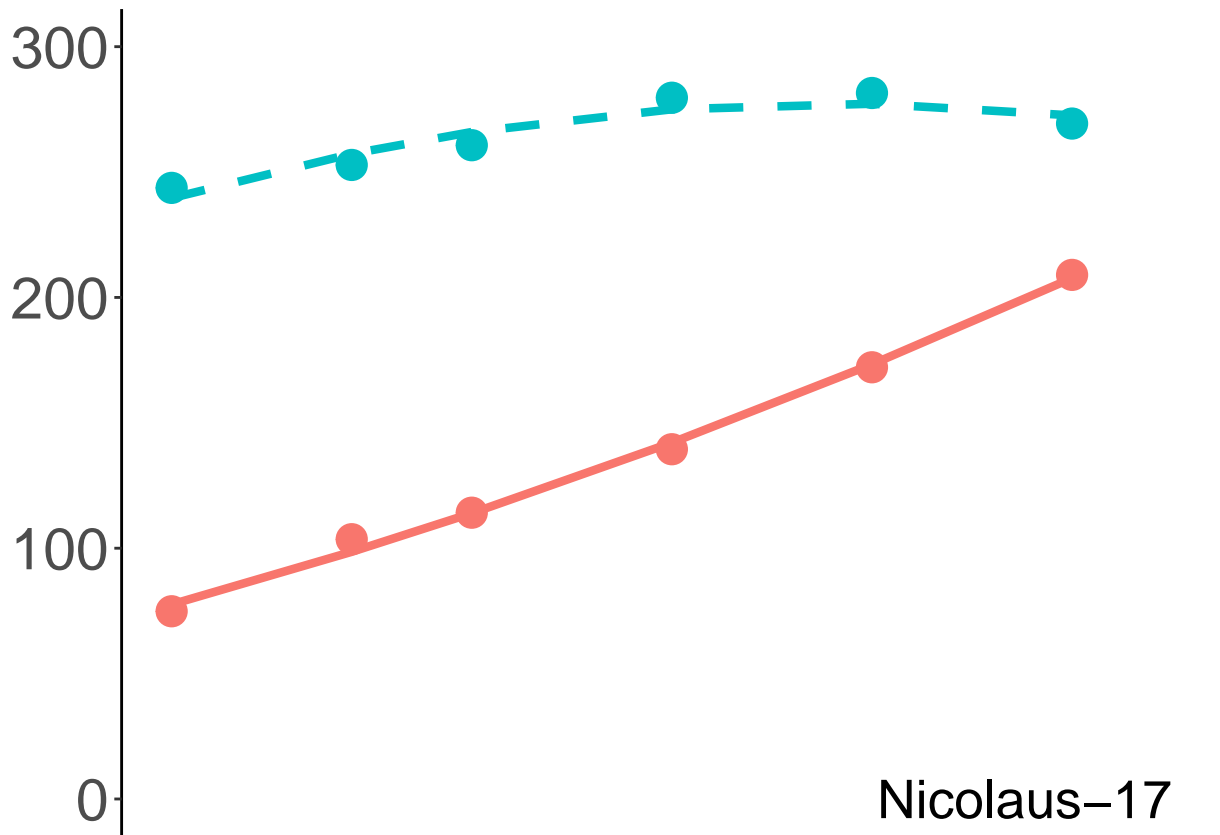
nic17_plot <- ggplot(data = suppl_data1_nic17 , aes( x = N_level_kgha , y = PI_N_Uptake)) +
  geom_point(data = suppl_data1_nic17 , aes ( x = N_level_kgha , y = PI_N_Uptake) , color = "#F8766D" ,
  geom_point(data = suppl_data1_nic17 , aes ( x = N_level_kgha , y = GrainYield_Mgha*23) , color = "#00BFC4" ,
  geom_line(data = nic17_df , aes( x = suppl_data1_nic17.N_level_kgha , y = nic17_fitted) , color = "#F8766D" ,
  geom_line(data = nic17_df_2 , aes( x = suppl_data1_nic17.N_level_kgha , y = nic17_fitted_2) , color = "#00BFC4" ,
  scale_y_continuous(breaks = seq(0 , 300 , by = 100) , sec.axis = sec_axis(~./23 , breaks = seq(0 , 10 , by = 1) ,
  coord_cartesian(ylim = c(0 , 300) , xlim = c(0 , 250)) +
  scale_x_continuous(breaks = seq(0 , 280 , by = 100)) +
  theme_classic() +
  theme(axis.text.y = element_text(size = 22),
```

```

axis.text.y.right = element_blank(),
axis.text.x = element_blank()) +
annotate("text" , x = 250 , y = 0 , label = "Nicolaus-17" , size = 7 , hjust = 1) +
labs( x = NULL , y = NULL)

```

nic17_plot



Williams-17

```

suppl_data1_wil17 <- suppl_data1 %>%
  filter(site_year == "Williams-17")

wil17_nuptake_lm <- lm(PI_N_Uptake ~ N_level_kgha + I(N_level_kgha*N_level_kgha) , data = suppl_data1_wil17)

wil17_fitted <- fitted(wil17_nuptake_lm)
wil17_df <- data.frame(suppl_data1_wil17$N_level_kgha , wil17_fitted) #creates dataframe

wil17_yield_lm <- lm(GrainYield_Mgha ~ N_level_kgha + I(N_level_kgha*N_level_kgha) , data = suppl_data1_wil17)

wil17_fitted_2 <- (fitted(wil17_yield_lm)*23)
wil17_df_2 <- data.frame(suppl_data1_wil17$N_level_kgha , wil17_fitted_2) #creates dataframe

wil17_plot <- ggplot(data = suppl_data1_wil17 , aes( x = N_level_kgha , y = PI_N_Uptake)) +
  geom_point(data = suppl_data1_wil17 , aes ( x = N_level_kgha , y = PI_N_Uptake) , color = "#F8766D" ,
  geom_point(data = suppl_data1_wil17 , aes ( x = N_level_kgha , y = GrainYield_Mgha*23) , color = "#00BFC4" ,

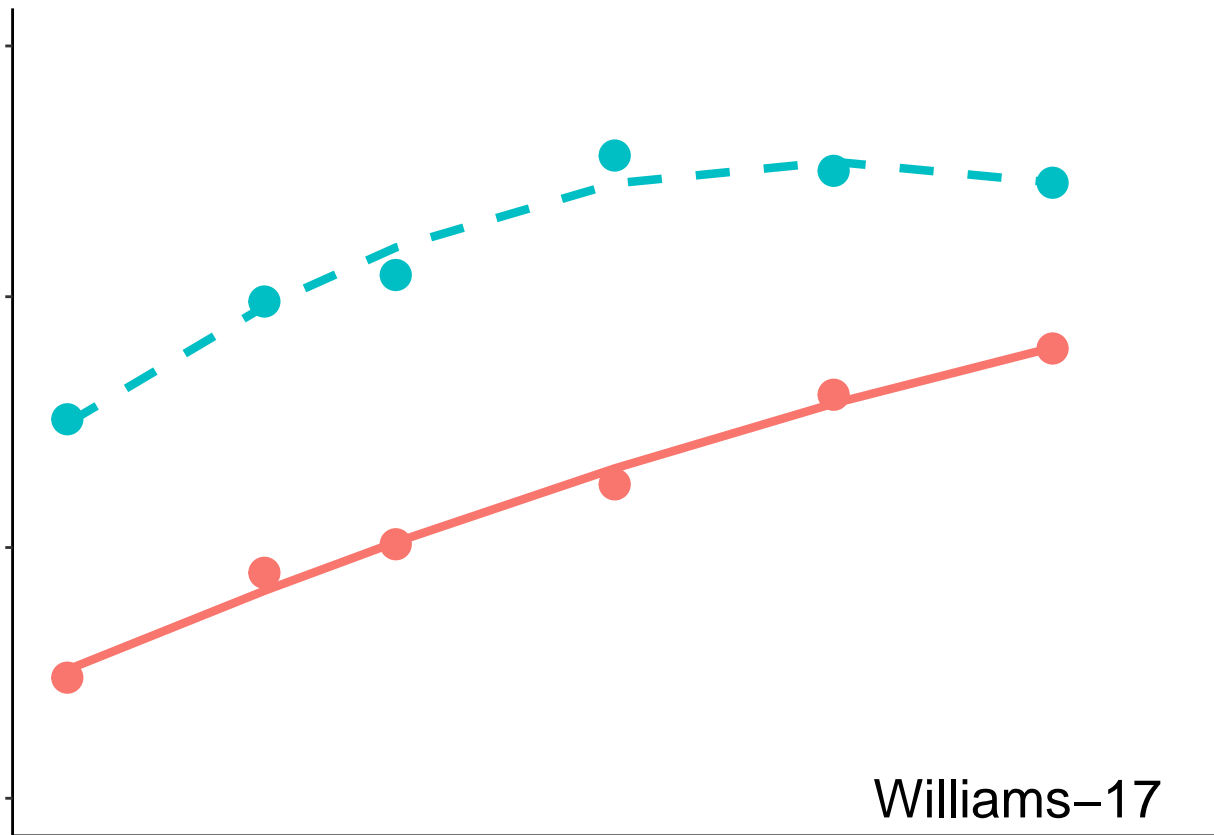
```

```

geom_line(data = wil17_df , aes( x = suppl_data1_wil17.N_level_kgha , y = wil17_fitted) , color = "#F
geom_line(data = wil17_df_2 , aes( x = suppl_data1_wil17.N_level_kgha , y = wil17_fitted_2) , color =
scale_y_continuous(breaks = seq(0 , 300 , by = 100) , sec.axis = sec_axis(~./23 , breaks = seq(0 , 1
coord_cartesian(ylim = c(0 , 300) , xlim = c(0 , 250)) +
scale_x_continuous(breaks = seq(0 , 280 , by = 100)) +
theme_classic() +
theme(axis.text.y = element_blank(),
      axis.text.x = element_blank()) +
annotate("text" , x = 250 , y = 0 , label = "Williams-17" , size = 7 , hjust = 1) +
labs( x = NULL , y = NULL)

```

wil17_plot



Arbuckle-18

```

suppl_data1_arb18 <- suppl_data1 %>%
  filter(site_year == "Arbuckle-18")

arb18_nuptake_lm <- lm(PI_N_Uptake ~ N_level_kgha + I(N_level_kgha*N_level_kgha) , data = suppl_data1_a

arb18_fitted <- fitted(arb18_nuptake_lm)
arb18_df <- data.frame(suppl_data1_arb18$N_level_kgha , arb18_fitted) #creates dataframe

arb18_yield_lm <- lm(GrainYield_Mgha ~ N_level_kgha + I(N_level_kgha*N_level_kgha) , data = suppl_data1

```



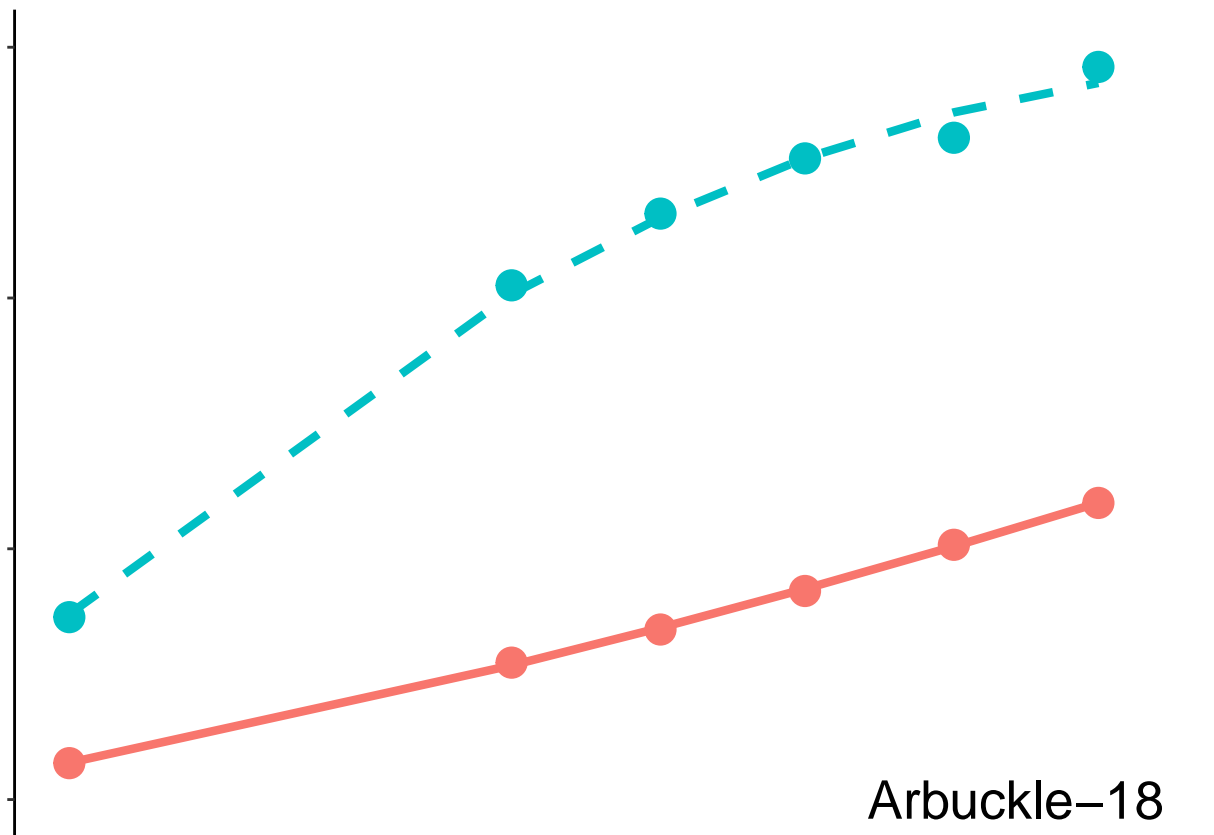
```

arb18_fitted_2 <- (fitted(arb18_yield_lm)*23)
arb18_df_2 <- data.frame(suppl_data1_arb18$N_level_kgha , arb18_fitted_2) #creates dataframe

arb18_plot <- ggplot(data = suppl_data1_arb18 , aes( x = N_level_kgha , y = PI_N_Uptake)) +
  geom_point(data = suppl_data1_arb18 , aes ( x = N_level_kgha , y = PI_N_Uptake) , color = "#F8766D") +
  geom_point(data = suppl_data1_arb18 , aes ( x = N_level_kgha , y = GrainYield_Mgha*23) , color = "#00BFC4") +
  geom_line(data = arb18_df , aes( x = suppl_data1_arb18.N_level_kgha , y = arb18_fitted) , color = "#F8766D") +
  geom_line(data = arb18_df_2 , aes( x = suppl_data1_arb18.N_level_kgha , y = arb18_fitted_2) , color = "#00BFC4") +
  scale_y_continuous(breaks = seq(0 , 300 , by = 100) , sec.axis = sec_axis(~./23 , breaks = seq(0 , 10 , by = 1))) +
  coord_cartesian(ylim = c(0 , 300) , xlim = c(0 , 250)) +
  scale_x_continuous(breaks = seq(0 , 280 , by = 100)) +
  theme_classic() +
  theme(axis.text.y = element_blank(),
        axis.text.x = element_blank()) +
  annotate("text" , x = 250 , y = 0 , label = "Arbuckle-18" , size = 7 , hjust = 1) +
  labs( x = NULL , y = NULL)

```

arb18_plot



Biggs-18

```

suppl_data1_biggs18 <- suppl_data1 %>%
  filter(site_year == "Biggs-18" )

biggs18_nuptake_lm <- lm(PI_N_Uptake ~ N_level_kgha + I(N_level_kgha*N_level_kgha) , data = suppl_data1.

```

```

biggs18_fitted <- fitted(biggs18_nuptake_lm)
biggs18_df <- data.frame(suppl_data1_biggs18$N_level_kgha , biggs18_fitted) #creates dataframe

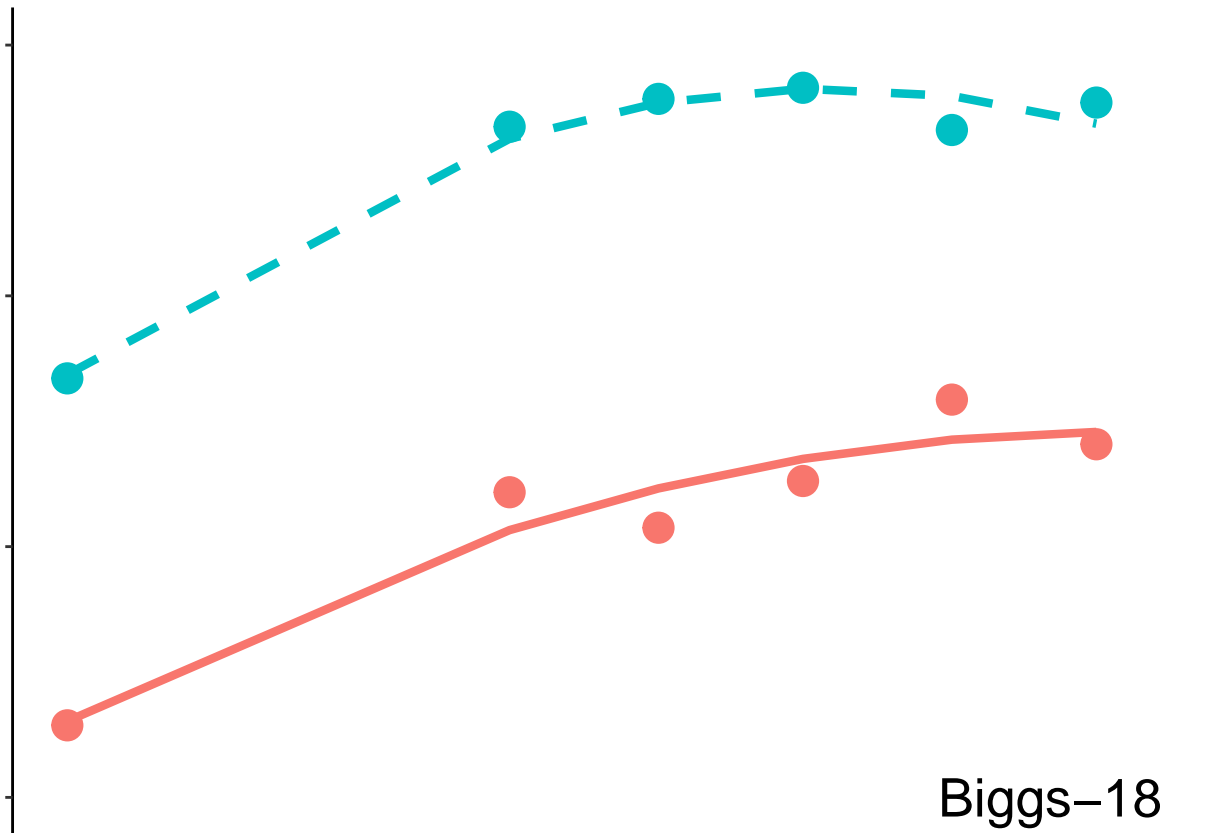
biggs18_yield_lm <- lm(GrainYield_Mgha ~ N_level_kgha + I(N_level_kgha*N_level_kgha) , data = suppl_data1_biggs18)

biggs18_fitted_2 <- (fitted(biggs18_yield_lm)*23)
biggs18_df_2 <- data.frame(suppl_data1_biggs18$N_level_kgha , biggs18_fitted_2) #creates dataframe

biggs18_plot <- ggplot(data = suppl_data1_biggs18 , aes( x = N_level_kgha , y = PI_N_Uptake)) +
  geom_point(data = suppl_data1_biggs18 , aes ( x = N_level_kgha , y = PI_N_Uptake) , color = "#F8766D") +
  geom_point(data = suppl_data1_biggs18 , aes ( x = N_level_kgha , y = GrainYield_Mgha*23) , color = "#F8766D") +
  geom_line(data = biggs18_df , aes( x = suppl_data1_biggs18.N_level_kgha , y = biggs18_fitted) , color = "#F8766D") +
  geom_line(data = biggs18_df_2 , aes( x = suppl_data1_biggs18.N_level_kgha , y = biggs18_fitted_2) , color = "#F8766D") +
  scale_y_continuous(breaks = seq(0 , 300 , by = 100) , sec.axis = sec_axis(~./23 , breaks = seq(0 , 10 , by = 1))) +
  coord_cartesian(ylim = c(0 , 300) , xlim = c(0 , 250)) +
  scale_x_continuous(breaks = seq(0 , 280 , by = 100)) +
  theme_classic() +
  theme(axis.text.y = element_blank(),
        axis.text.x = element_blank()) +
  annotate("text" , x = 250 , y = 0 , label = "Biggs-18" , size = 7 , hjust = 1) +
  labs( x = NULL , y = NULL)

```

biggs18_plot



Marysville-18

```
suppl_data1_mry18 <- suppl_data1 %>%
  filter(site_year == "Marysville-18" )

mry18_nuptake_lm <- lm(PI_N_Uptake ~ N_level_kgha + I(N_level_kgha*N_level_kgha) , data = suppl_data1_mry18)

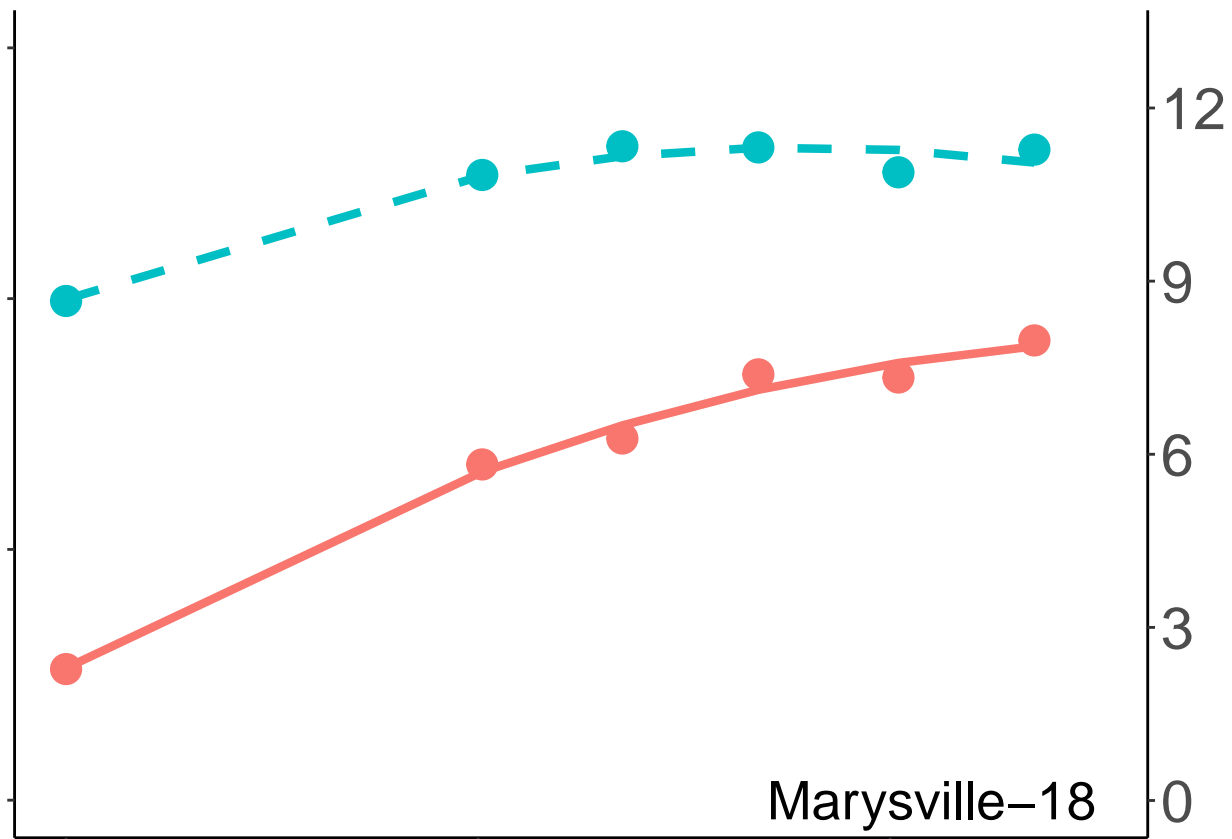
mry18_fitted <- fitted(mry18_nuptake_lm)
mry18_df <- data.frame(suppl_data1_mry18$N_level_kgha , mry18_fitted) #creates dataframe

mry18_yield_lm <- lm(GrainYield_Mgha ~ N_level_kgha + I(N_level_kgha*N_level_kgha) , data = suppl_data1_mry18)

mry18_fitted_2 <- (fitted(mry18_yield_lm)*23)
mry18_df_2 <- data.frame(suppl_data1_mry18$N_level_kgha , mry18_fitted_2) #creates dataframe

mry18_plot <- ggplot(data = suppl_data1_mry18 , aes( x = N_level_kgha , y = PI_N_Uptake)) +
  geom_point(data = suppl_data1_mry18 , aes ( x = N_level_kgha , y = PI_N_Uptake) , color = "#F8766D" ,
  geom_point(data = suppl_data1_mry18 , aes ( x = N_level_kgha , y = GrainYield_Mgha*23) , color = "#0072B2" ,
  geom_line(data = mry18_df , aes( x = suppl_data1_mry18.N_level_kgha , y = mry18_fitted) , color = "#F8766D" ,
  geom_line(data = mry18_df_2 , aes( x = suppl_data1_mry18.N_level_kgha , y = mry18_fitted_2) , color = "#0072B2" ,
  scale_y_continuous(breaks = seq(0 , 300 , by = 100) , sec.axis = sec_axis(~./23 , breaks = seq(0 , 10) , labels = seq(0 , 10) ,
  coord_cartesian(ylim = c(0 , 300) , xlim = c(0 , 250)) +
  scale_x_continuous(breaks = seq(0 , 280 , by = 100)) +
  theme_classic() +
  theme(axis.text.y.right = element_text(size = 22),
        axis.text.y.left = element_blank(),
        axis.text.x = element_blank()) +
  annotate("text" , x = 250 , y = 0 , label = "Marysville-18" , size = 7 , hjust = 1) +
  labs( x = NULL , y = NULL)

mry18_plot
```



Nicolaus-18

```

suppl_data1_nic18 <- suppl_data1 %>%
  filter(site_year == "Nicolaus-18" )

nic18_nuptake_lm <- lm(PI_N_Uptake ~ N_level_kgha + I(N_level_kgha*N_level_kgha) , data = suppl_data1_nic18)

nic18_fitted <- fitted(nic18_nuptake_lm)
nic18_df <- data.frame(suppl_data1_nic18$N_level_kgha , nic18_fitted) #creates dataframe

nic18_yield_lm <- lm(GrainYield_Mgha ~ N_level_kgha + I(N_level_kgha*N_level_kgha) , data = suppl_data1_nic18)

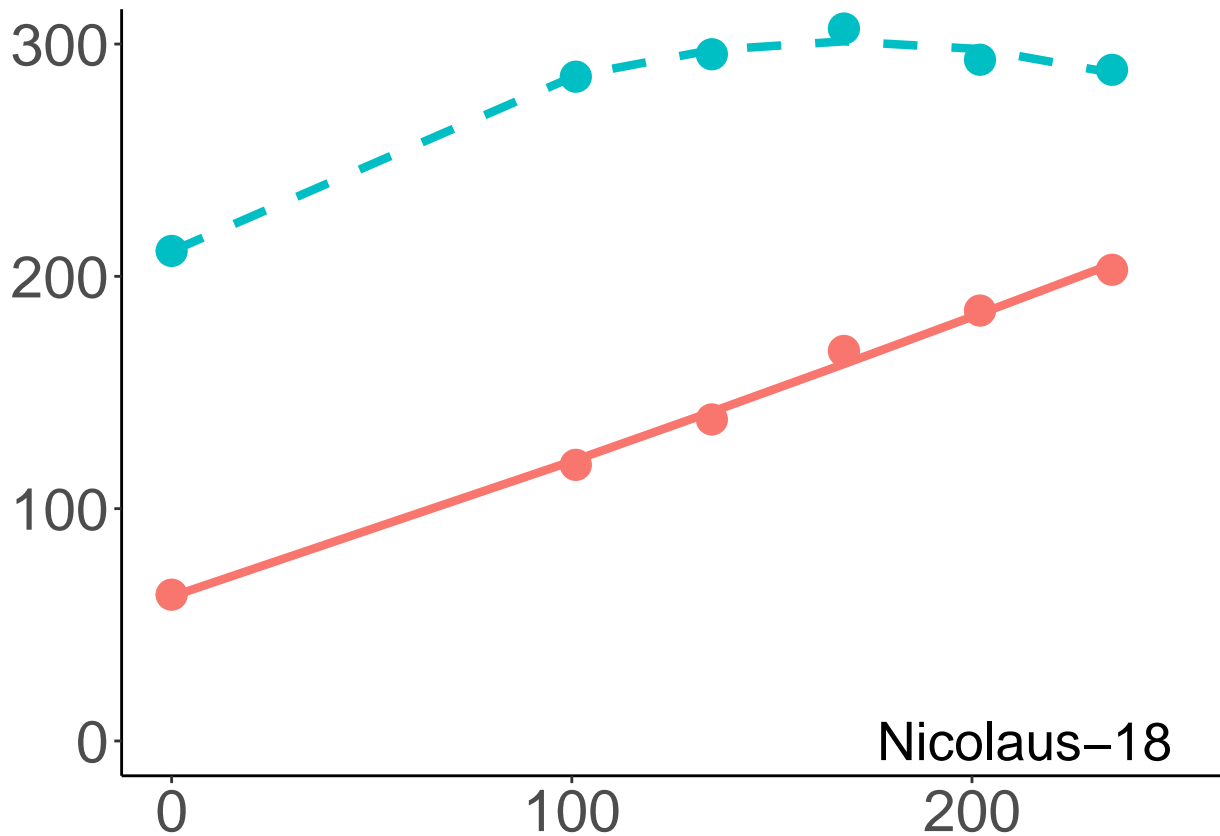
nic18_fitted_2 <- (fitted(nic18_yield_lm)*23)
nic18_df_2 <- data.frame(suppl_data1_nic18$N_level_kgha , nic18_fitted_2) #creates dataframe

nic18_plot <- ggplot(data = suppl_data1_nic18 , aes( x = N_level_kgha , y = PI_N_Uptake)) +
  geom_point(data = suppl_data1_nic18 , aes ( x = N_level_kgha , y = PI_N_Uptake) , color = "#F8766D" ,
  geom_point(data = suppl_data1_nic18 , aes ( x = N_level_kgha , y = GrainYield_Mgha*23) , color = "#0072B2" ,
  geom_line(data = nic18_df , aes( x = suppl_data1_nic18.N_level_kgha , y = nic18_fitted) , color = "#F8766D" ,
  geom_line(data = nic18_df_2 , aes( x = suppl_data1_nic18.N_level_kgha , y = nic18_fitted_2) , color = "#0072B2" ,
  scale_y_continuous(breaks = seq(0 , 300 , by = 100) , sec.axis = sec_axis(~./23 , breaks = seq(0 , 12 , by = 3)) ,
  coord_cartesian(ylim = c(0 , 300) , xlim = c(0 , 250)) +
  scale_x_continuous(breaks = seq(0 , 280 , by = 100)) +
  theme_classic() +
  theme(axis.text.y.right = element_blank(),
        axis.text.x = element_text(size = 22),

```

```
axis.text.y.left = element_text(size = 22)) +
  annotate("text" , x = 250 , y = 0 , label = "Nicolaus-18" , size = 7 , hjust = 1) +
  labs( x = NULL , y = NULL)
```

nic18_plot



Arbuckle-19

```
suppl_data1_arb19 <- suppl_data1 %>%
  filter(site_year == "Arbuckle-19" )

arb19_nuptake_lm <- lm(PI_N_Uptake ~ N_level_kgha + I(N_level_kgha*N_level_kgha) , data = suppl_data1_arb19)

arb19_fitted <- fitted(arb19_nuptake_lm)
arb19_df <- data.frame(suppl_data1_arb19$N_level_kgha , arb19_fitted) #creates dataframe

arb19_yield_lm <- lm(GrainYield_Mgha ~ N_level_kgha + I(N_level_kgha*N_level_kgha) , data = suppl_data1_arb19)

arb19_fitted_2 <- (fitted(arb19_yield_lm)*23)
arb19_df_2 <- data.frame(suppl_data1_arb19$N_level_kgha , arb19_fitted_2) #creates dataframe

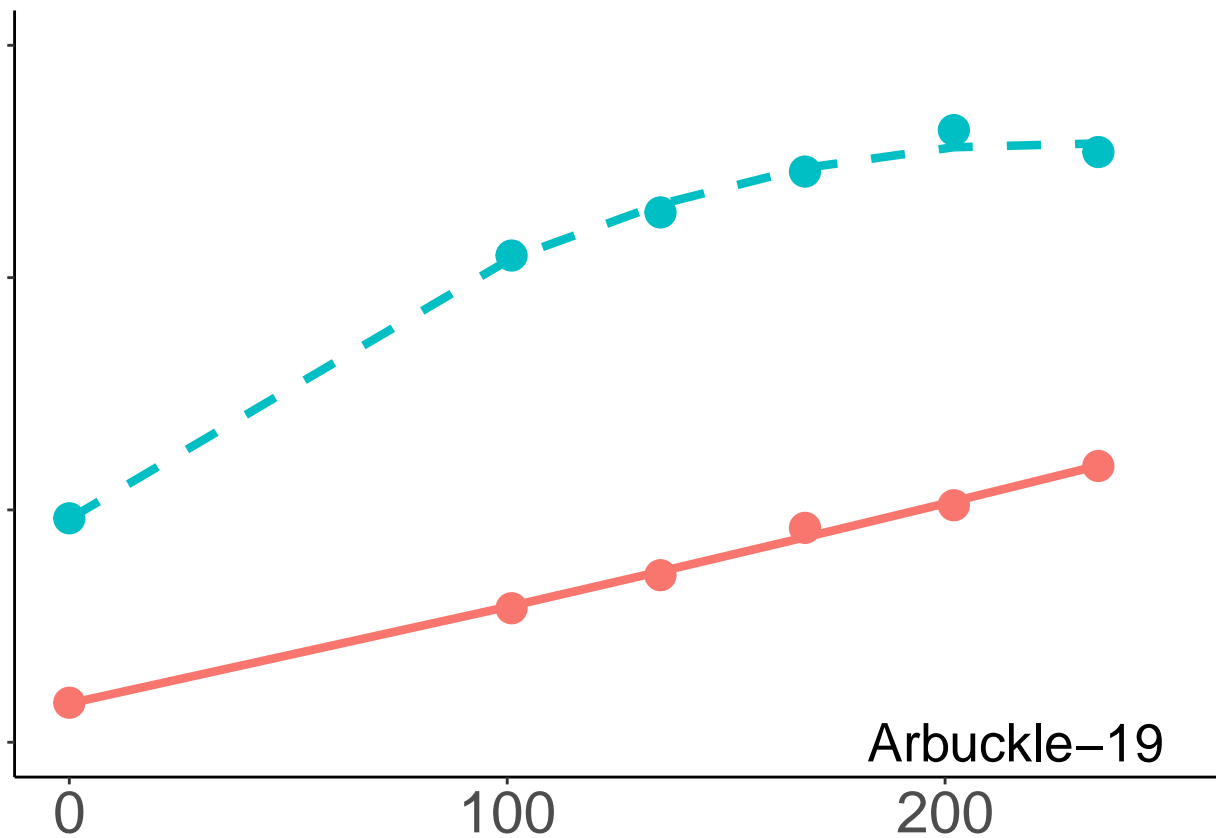
arb19_plot <- ggplot(data = suppl_data1_arb19 , aes( x = N_level_kgha , y = PI_N_Uptake)) +
  geom_point(data = suppl_data1_arb19 , aes ( x = N_level_kgha , y = PI_N_Uptake) , color = "#F8766D" ,
  geom_point(data = suppl_data1_arb19 , aes ( x = N_level_kgha , y = GrainYield_Mgha*23) , color = "#00BFC4" ,
  geom_line(data = arb19_df , aes( x = suppl_data1_arb19.N_level_kgha , y = arb19_fitted) , color = "#F8766D")
```

```

geom_line(data = arb19_df_2 , aes( x = suppl_data1_arb19.N_level_kgha , y = arb19_fitted_2) , color =
scale_y_continuous(breaks = seq(0 , 300 , by = 100) , sec.axis = sec_axis(~./23 , breaks = seq(0 , 1
coord_cartesian(ylim = c(0 , 300) , xlim = c(0 , 250)) +
scale_x_continuous(breaks = seq(0 , 280 , by = 100)) +
theme_classic() +
theme(axis.text.y = element_blank(),
      axis.text.x = element_text(size = 22)) +
annotate("text" , x = 250 , y = 0 , label = "Arbuckle-19" , size = 7 , hjust = 1) +
labs( x = NULL , y = NULL)

```

arb19_plot



Davis-19

```

suppl_data1_davis19 <- suppl_data1 %>%
  filter(site_year == "Davis-19" )

davis19_nuptake_lm <- lm(PI_N_Uptake ~ N_level_kgha + I(N_level_kgha*N_level_kgha) , data = suppl_data1
davis19_fitted <- fitted(davis19_nuptake_lm)
davis19_df <- data.frame(suppl_data1_davis19$N_level_kgha , davis19_fitted) #creates dataframe

davis19_yield_lm <- lm(GrainYield_Mgha ~ N_level_kgha + I(N_level_kgha*N_level_kgha) , data = suppl_data
davis19_fitted_2 <- (fitted(davis19_yield_lm)*23)

```

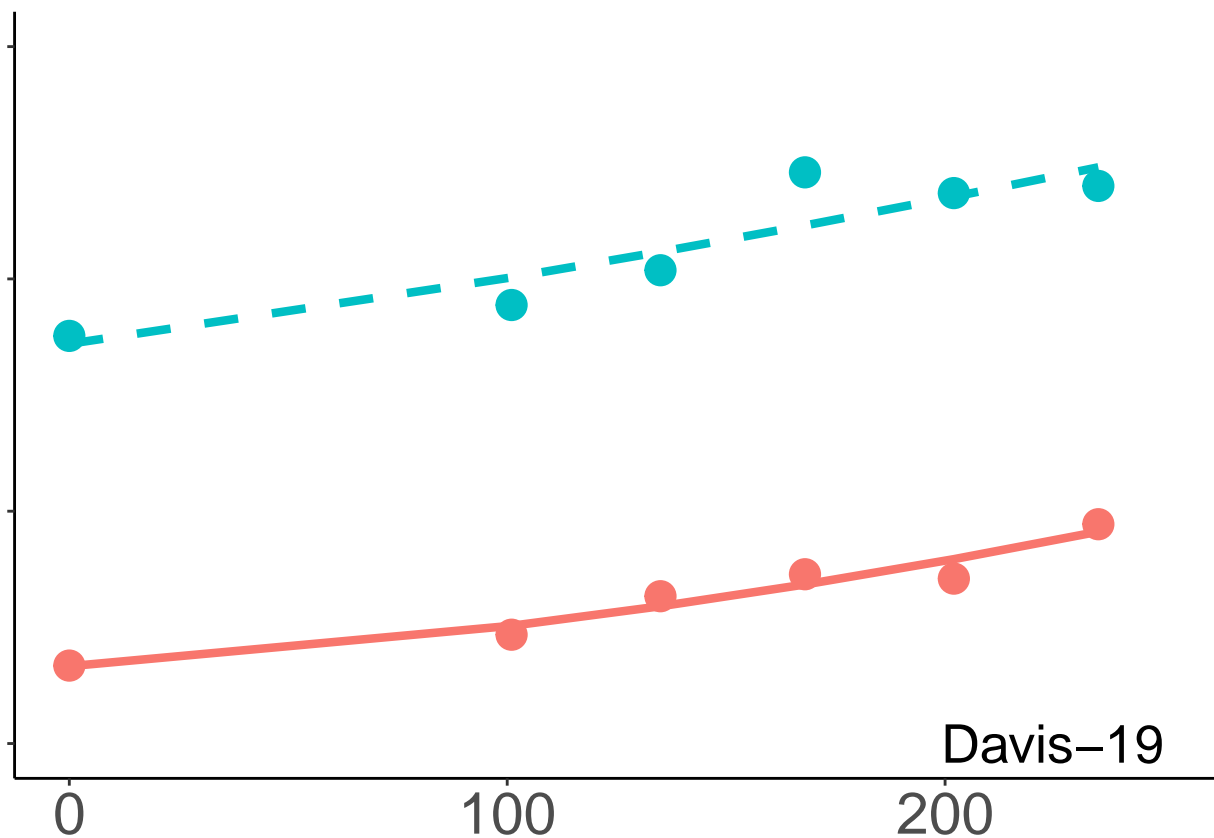
```

davis19_df_2 <- data.frame(suppl_data1_davis19$N_level_kgha , davis19_fitted_2) #creates dataframe

davis19_plot <- ggplot(data = suppl_data1_davis19 , aes( x = N_level_kgha , y = PI_N_Uptake)) +
  geom_point(data = suppl_data1_davis19 , aes ( x = N_level_kgha , y = PI_N_Uptake) , color = "#F8766D") +
  geom_point(data = suppl_data1_davis19 , aes ( x = N_level_kgha , y = GrainYield_Mgha*23) , color = "#F8766D") +
  geom_line(data = davis19_df , aes( x = suppl_data1_davis19.N_level_kgha , y = davis19_fitted) , color = "#F8766D") +
  geom_line(data = davis19_df_2 , aes( x = suppl_data1_davis19.N_level_kgha , y = davis19_fitted_2) , color = "#F8766D") +
  scale_y_continuous(breaks = seq(0 , 300 , by = 100) , sec.axis = sec_axis(~./23 , breaks = seq(0 , 10 , by = 1))) +
  coord_cartesian(ylim = c(0 , 300) , xlim = c(0 , 250)) +
  scale_x_continuous(breaks = seq(0 , 280 , by = 100)) +
  theme_classic() +
  theme(axis.text.y = element_blank(),
        axis.text.x = element_text(size = 22)) +
  annotate("text" , x = 250 , y = 0 , label = "Davis-19" , size = 7 , hjust = 1) +
  labs( x = NULL , y = NULL)

```

davis19_plot



Marysville-19

```

suppl_data1_mry19 <- suppl_data1 %>%
  filter(site_year == "Marysville-19" )

mry19_nuptake_lm <- lm(PI_N_Uptake ~ N_level_kgha + I(N_level_kgha*N_level_kgha) , data = suppl_data1_mry19)

```

```

mry19_fitted <- fitted(mry19_nuptake_lm)
mry19_df <- data.frame(suppl_data1_mry19$N_level_kgha , mry19_fitted) #creates dataframe

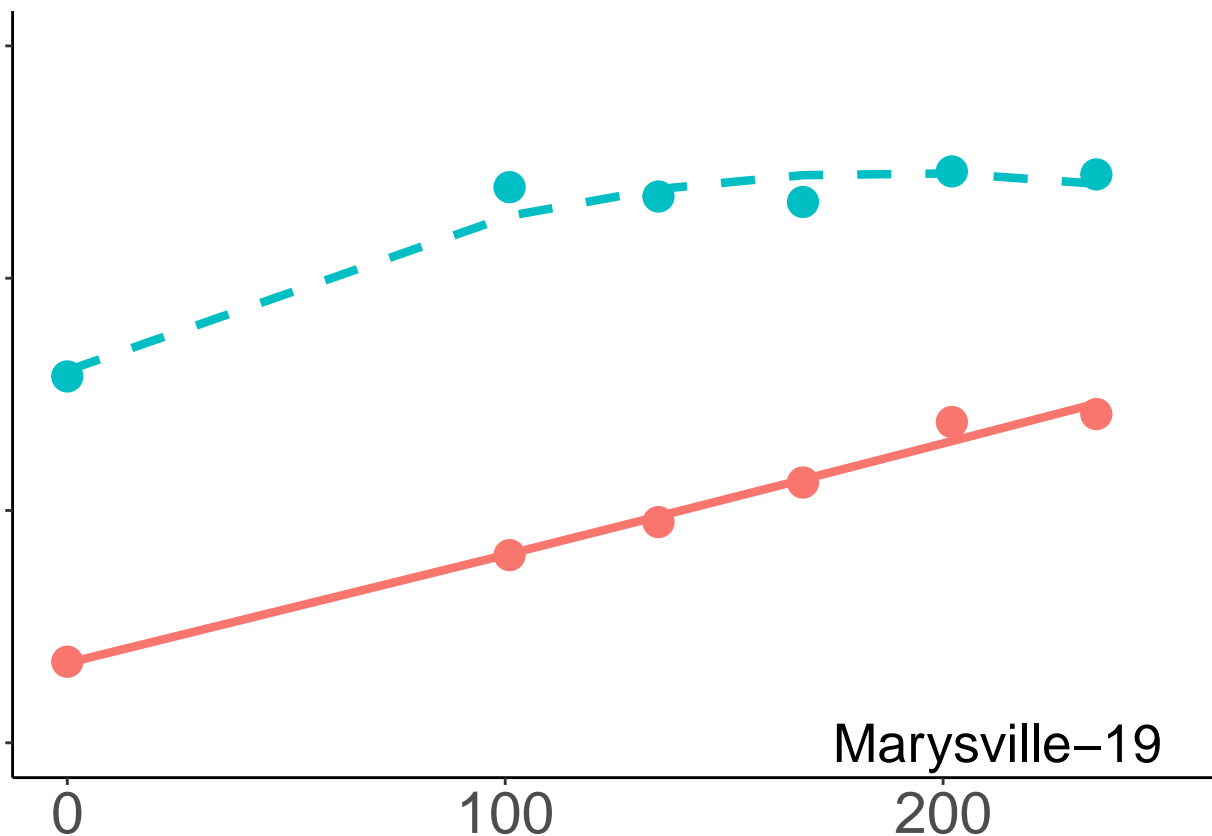
mry19_yield_lm <- lm(GrainYield_Mgha ~ N_level_kgha + I(N_level_kgha*N_level_kgha) , data = suppl_data1

mry19_fitted_2 <- (fitted(mry19_yield_lm)*23)
mry19_df_2 <- data.frame(suppl_data1_mry19$N_level_kgha , mry19_fitted_2) #creates dataframe

mry19_plot <- ggplot(data = suppl_data1_mry19 , aes( x = N_level_kgha , y = PI_N_Uptake)) +
  geom_point(data = suppl_data1_mry19 , aes ( x = N_level_kgha , y = PI_N_Uptake) , color = "#F8766D" ,
  geom_point(data = suppl_data1_mry19 , aes ( x = N_level_kgha , y = GrainYield_Mgha*23) , color = "#00BFC4" ,
  geom_line(data = mry19_df , aes( x = suppl_data1_mry19.N_level_kgha , y = mry19_fitted) , color = "#F8766D" ,
  geom_line(data = mry19_df_2 , aes( x = suppl_data1_mry19.N_level_kgha , y = mry19_fitted_2) , color = "#00BFC4" ,
  scale_y_continuous(breaks = seq(0 , 300 , by = 100) , sec.axis = sec_axis(~./23 , breaks = seq(0 , 10 , by = 1) ,
  coord_cartesian(ylim = c(0 , 300) , xlim = c(0 , 250)) +
  scale_x_continuous(breaks = seq(0 , 280 , by = 100)) +
  theme_classic() +
  theme(axis.text.x = element_text(size = 22),
        axis.text.y = element_blank()) +
  annotate("text" , x = 250 , y = 0 , label = "Marysville-19" , size = 7 , hjust = 1) +
  labs( x = NULL , y = NULL)

```

mry19_plot



RES-19

```
suppl_data1_res19 <- suppl_data1 %>%
  filter(site_year == "RES-19" )

res19_nuptake_lm <- lm(PI_N_Uptake ~ N_level_kgha + I(N_level_kgha*N_level_kgha) , data = suppl_data1_r

res19_fitted <- fitted(res19_nuptake_lm)
res19_df <- data.frame(suppl_data1_res19$N_level_kgha , res19_fitted) #creates dataframe

res19_yield_lm <- lm(GrainYield_Mgha ~ N_level_kgha + I(N_level_kgha*N_level_kgha) , data = suppl_data1.

res19_fitted_2 <- (fitted(res19_yield_lm)*23)
res19_df_2 <- data.frame(suppl_data1_res19$N_level_kgha , res19_fitted_2) #creates dataframe

res19_plot <- ggplot(data = suppl_data1_res19 , aes( x = N_level_kgha , y = PI_N_Uptake)) +
  geom_point(data = suppl_data1_res19 , aes ( x = N_level_kgha , y = PI_N_Uptake) , color = "#F8766D" ,
  geom_point(data = suppl_data1_res19 , aes ( x = N_level_kgha , y = GrainYield_Mgha*23) , color = "#00
  geom_line(data = res19_df , aes( x = suppl_data1_res19.N_level_kgha , y = res19_fitted) , color = "#F
  geom_line(data = res19_df_2 , aes( x = suppl_data1_res19.N_level_kgha , y = res19_fitted_2) , color =
  scale_y_continuous(breaks = seq(0 , 300 , by = 100) , sec.axis = sec_axis(~./23 , breaks = seq(0 , 1
  coord_cartesian(ylim = c(0 , 300) , xlim = c(0 , 250)) +
  scale_x_continuous(breaks = seq(0 , 280 , by = 100)) +
  theme_classic() +
  theme(axis.text.y.left = element_blank(),
        axis.text.x = element_text(size = 22),
        axis.text.y.right = element_text(size = 22)) +
  annotate("text" , x = 250 , y = 0 , label = "RES-19" , size = 7 , hjust = 1) +
  labs( x = NULL , y = NULL)

res19_plot
```

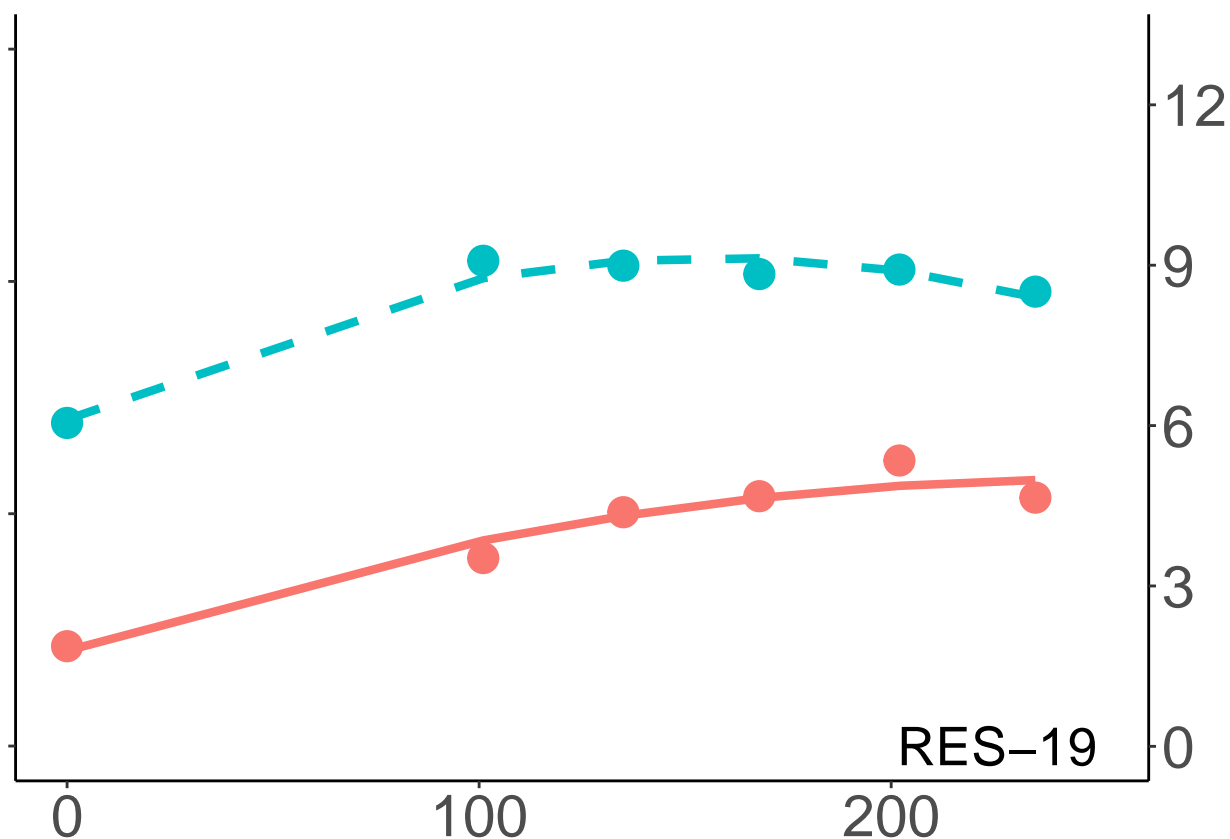
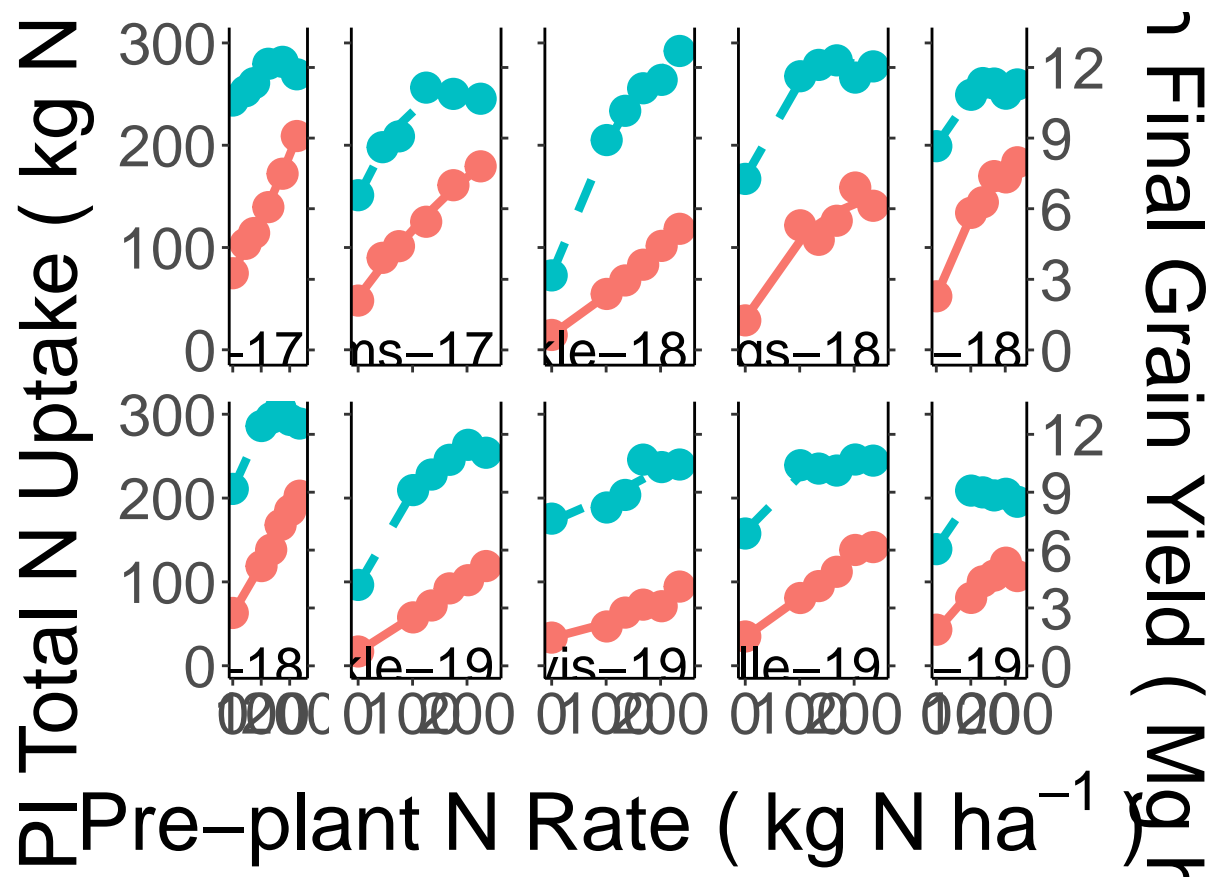
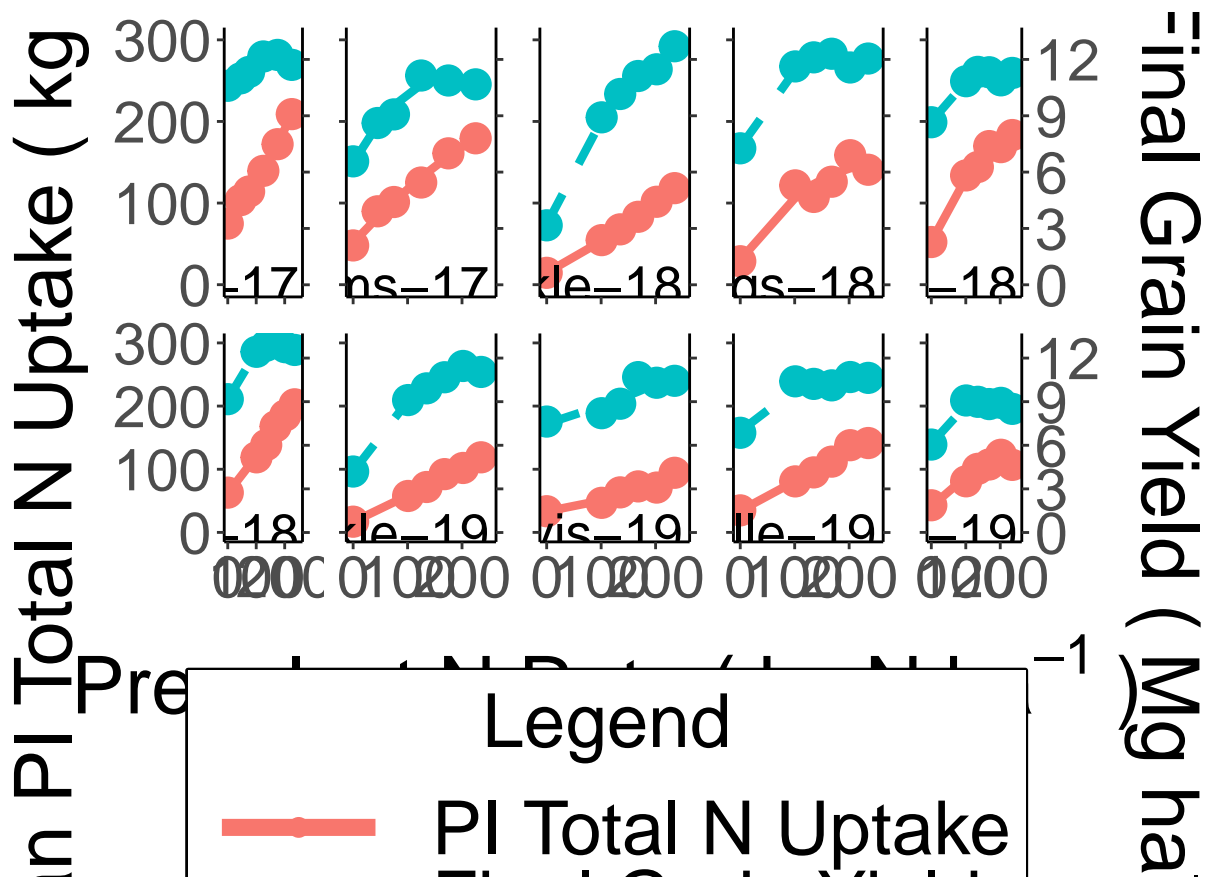


FIGURE 2

```
Fig2 <- grid.arrange(arrangeGrob(
  nic17_plot,
  wil17_plot,
  arb18_plot,
  biggs18_plot,
  mry18_plot,
  nic18_plot,
  arb19_plot,
  davis19_plot,
  mry19_plot,
  res19_plot,
  ncol = 5,
  nrow = 2 ,
  widths = c(1.4, 1.2 , 1.2, 1.2, 1.3 ),
  bottom = textGrob("Pre-plant N Rate ( kg N ha-1~") ,
    gp = gpar( fontsize = 30)),
  right = textGrob("Mean Final Grain Yield ( Mg ha-1~") ,
    gp = gpar(fontsize = 30) ,
    rot = 270),
  left = textGrob("Mean PI Total N Uptake ( kg N ha-1~") ,
    gp = gpar( fontsize = 30) ,
    rot = 90)
))
```



```
Fig2 <- grid.arrange(arrangeGrob(Fig2,
  legend,
  ncol = 1,
  nrow = 2,
  heights = c(8 , 1.5)))
```



```
ggsave("FIGURES/Fig2.tiff" , Fig2, compression = "lzw" , width = 22 , height = 12 , type = "cairo" , dp
```

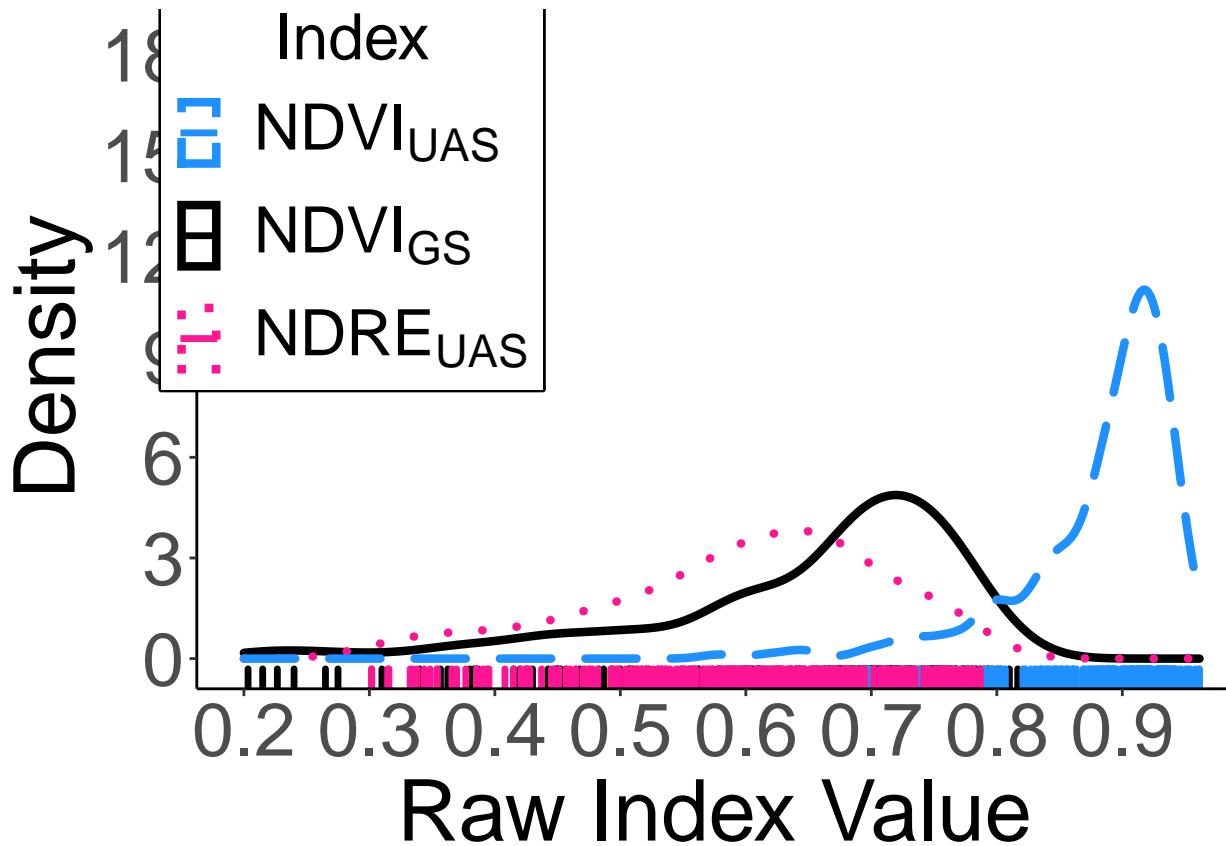
FIGURE 3

IV Kernel

```
Fig3.1 <- ggplot(data = paper3_data , aes(x = Index) ) +
  geom_density(data = paper3_data, aes(x = Index , color = Platform , linetype = Platform) , size = 1.1) +
  geom_rug(data = paper3_data , aes(x = Index , color = Platform) , size = 1.2) +
  theme_classic() +
  theme(axis.title = element_text(size = 32),
        axis.text = element_text(size = 28),
        legend.text = element_text(size = 24),
        legend.title = element_text(size = 24 , hjust = 0.5),
        legend.position = c(.15 , .75),
        legend.text.align = 0,
        legend.box.background = element_rect(size = 1)) +
  coord_cartesian(xlim = c(0.2 , 0.95) ,ylim = c(0 , 18)) +
  scale_x_continuous(breaks = seq(.2 , .9 , by = .1)) +
  scale_y_continuous(breaks = seq(0 , 18 , by = 3)) +
  labs(x = "Raw Index Value" , y = "Density" , color = "Index" , linetype = "Index") +
  scale_color_manual(breaks = c("sUAS NDVI" , "GreenSeeker NDVI" , "sUAS NDRE") , values = c( "dodgerblue" , "green" , "red" )) +
  scale_linetype_manual(breaks = c("sUAS NDVI" , "GreenSeeker NDVI" , "sUAS NDRE") , values = c("longdashed" , "solid" , "dotted" )) +
  guides(
    color = guide_legend(byrow = TRUE)
  ) +
```

```
theme(
  legend.spacing.y = unit(0.4, "cm")) +
  annotate("text" , x = (0.2) , y = (18) , label = "(a)" , size = 10 , hjust = 0.5 , fontface = 2)
```

Fig3.1



```
ggsave("FIGURES/Fig3.1.tiff" , Fig3.1 , compression = "lzw" , width = 15 , height = 10, type = "cairo"
```

SI Kernel

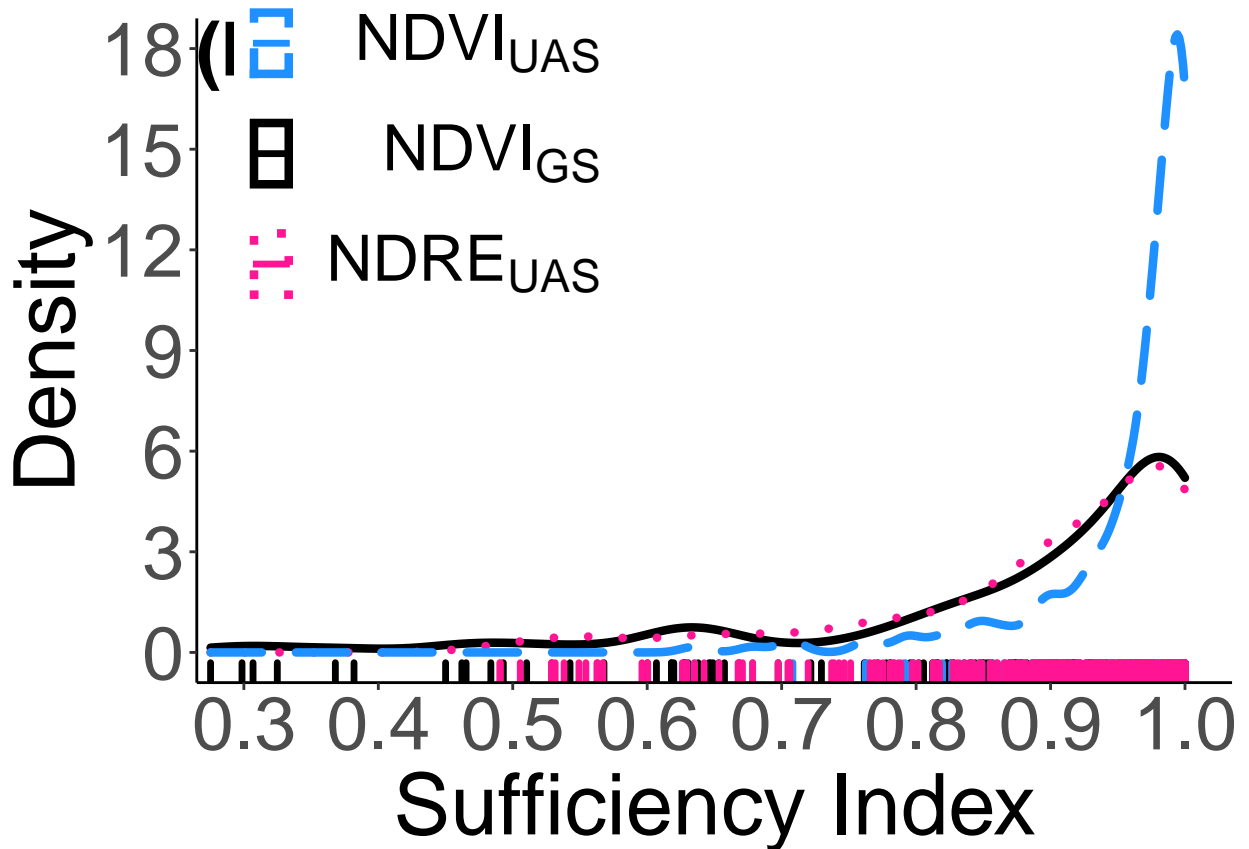
```
Fig3.2 <- ggplot(data = paper3_data , aes(x = SI) ) +
  geom_density(data = paper3_data, aes(x = SI , color = Platform , linetype = Platform) , size = 1.5) +
  geom_rug(data = paper3_data , aes(x = SI , color = Platform) , size = 1.2) +
  theme_classic() +
  theme(axis.title = element_text(size = 32),
        axis.text = element_text(size = 28),
        legend.text = element_text(size = 24),
        legend.title = element_text(size = 24),
        legend.position = c(0.22 , 0.87)
  ) +
  coord_cartesian(xlim = c(.3 , 1) , ylim = c(0 , 18)) +
  scale_x_continuous(breaks = seq(.3 , 1 , by = .1)) +
  scale_y_continuous(breaks = seq(0 , 18 , by = 3)) +
  labs(x = "Sufficiency Index" , y = "Density" , color = "Index" , linetype = "Index") +
  scale_color_manual(breaks = c("sUAS NDVI" , "GreenSeeker NDVI" , "sUAS NDRE") , values = c( "dodgerblue", "black", "pink"))
```

```

scale_linetype_manual(breaks = c("sUAS NDVI" , "GreenSeeker NDVI" , "sUAS NDRE") , values = c("longdashed", "solid", "dotted")) +
  guides(
    color = guide_legend(byrow = TRUE)
  ) +
  theme(
    legend.spacing.y = unit(0.5, "cm")) +
  annotate("text" , x = (0.3) , y = (18) , label = "(b)" , size = 10 , hjust = 0.5 , fontface = 2)

```

Fig3.2



```

ggsave("FIGURES/Fig3.2.tiff" , Fig3.2 , compression = "lzw" , width = 15 , height = 10, type = "cairo")

```

```

Fig3.2.2 <- Fig3.2 +
  labs(x = "Sufficiency Index" , y = NULL , color = "Index" , linetype = "Index") +
  theme(axis.title = element_text(size = 32),
        axis.text = element_text(size = 28),
        legend.text = element_text(size = 24),
        legend.title = element_text(size = 24),
        legend.position = "none"
  )

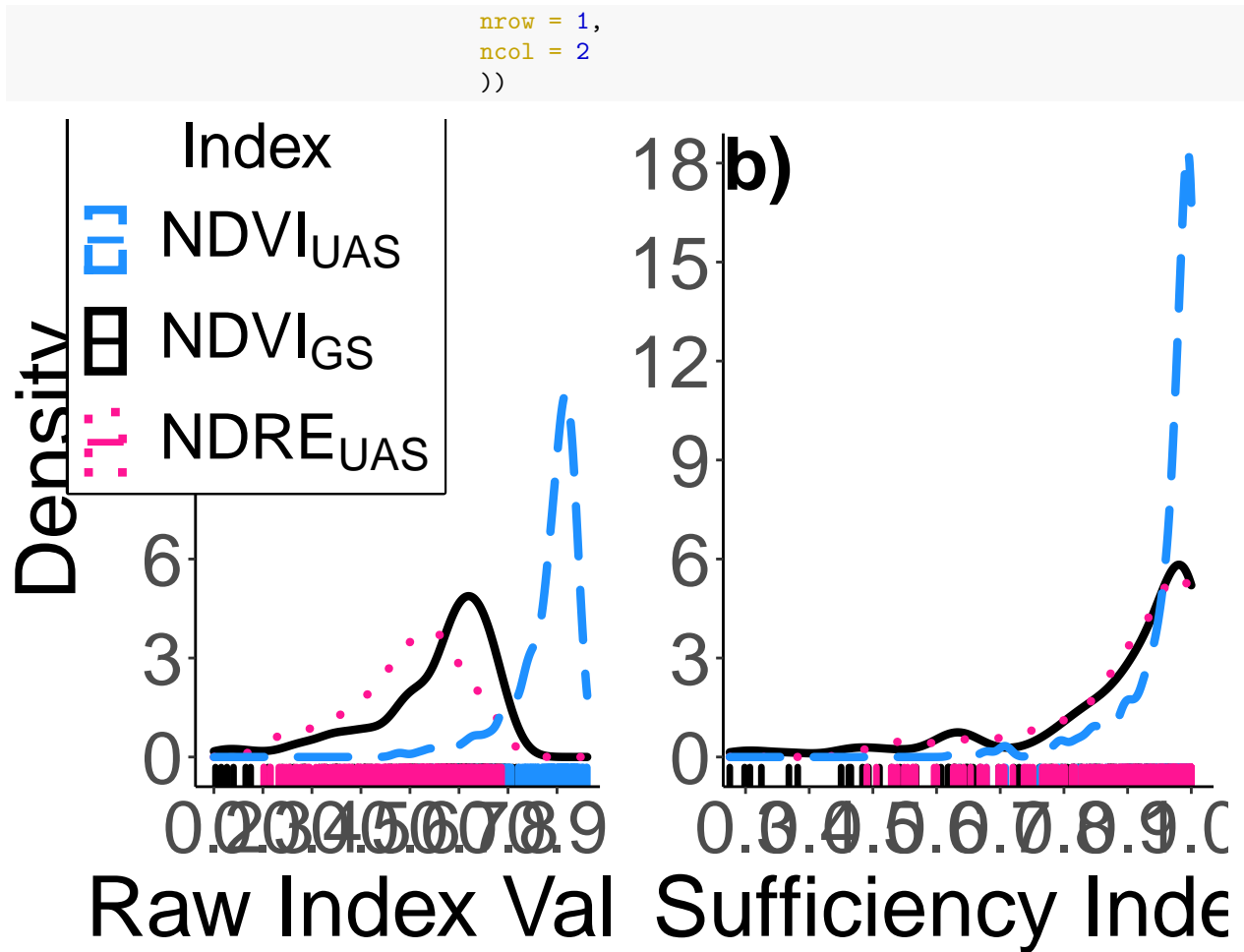
```

FIGURE 3

```

Fig3 <- grid.arrange(arrangeGrob(Fig3.1,
                                  Fig3.2.2,

```



```
ggsave("FIGURES/Fig3.tiff" , Fig3 , compression = "lzw" , width = 20 , height = 10, type = "cairo" , dp
```

FIGURE 4

gs-ndvi-SI

```
fit.lm <- lm(SI ~ PI_N_Uptake, data= paper3_gsdata)

a.ini <- fit.lm$coefficients[1]
b.ini <- fit.lm$coefficients[2]
clx.ini <- mean(paper3_gsdata$PI_N_Uptake)

quadplat <- function(x, a, b, clx) {
  ifelse(x < clx, a + b * x + (-0.5*b/clx) * x * x,
        a + b * clx + (-0.5*b/clx) * clx * clx)}

model <- nls(SI ~ quadplat(PI_N_Uptake, a, b, clx),
  data = paper3_gsdata,
  start = list(a = a.ini,
               b = b.ini,
               clx = clx.ini),
  trace = FALSE,
```

```

      nls.control(maxiter = 1000))

summary(model)

##
## Formula: SI ~ quadplat(PI_N_Uptake, a, b, clx)
##
## Parameters:
##      Estimate Std. Error t value Pr(>|t|)
## a    2.197e-01  3.757e-02   5.848 1.66e-08 ***
## b    1.318e-02  1.103e-03  11.946 < 2e-16 ***
## clx  1.126e+02  4.922e+00  22.868 < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.08152 on 234 degrees of freedom
##
## Number of iterations to convergence: 9
## Achieved convergence tolerance: 4.188e-06

nullfunct <- function(x, m){m}

m.ini      <- mean(paper3_gsdata$SI)

null <- nls(SI ~ nullfunct(PI_N_Uptake, m),
            data = paper3_gsdata,
            start = list(m = m.ini),
            trace = FALSE,
            nls.control(maxiter = 1000))

nagelkerke(model,
            null)

## $Models
##
## Model: "nls, SI ~ quadplat(PI_N_Uptake, a, b, clx), paper3_gsdata, list(a = a.ini, b = b.ini, clx = c
## Null:  "nls, SI ~ nullfunct(PI_N_Uptake, m), paper3_gsdata, list(m = m.ini), list(1000, 1e-05, 0.000
##
## $Pseudo.R.squared.for.model.vs.null
##                                Pseudo.R.squared
## McFadden                      -1.478860
## Cox and Snell (ML)             0.729047
## Nagelkerke (Cragg and Uhler)   -0.514101
##
## $Likelihood.ratio.test
## Df.diff LogLik.diff Chisq    p.value
##      -2      -154.74 309.48 6.2806e-68
##
## $Number.of.observations
##
## Model: 237
## Null:  237
##
## $Messages

```



```
## [1] "Note: For models fit with REML, these statistics are based on refitting with ML"
##
## $Warnings
## [1] "None"

confint2(model,
          level = 0.95)

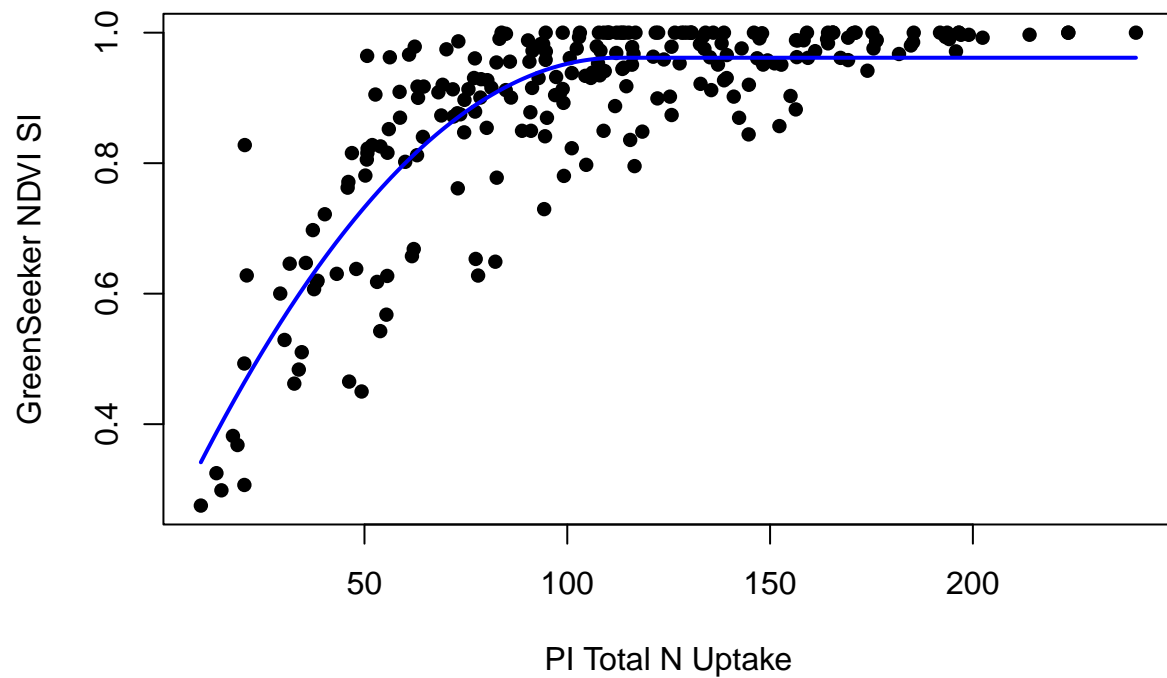
##           2.5 %           97.5 %
## a      0.14569776    0.29373689
## b      0.01100811    0.01535621
## clx 102.86320065 122.25856538

Boot <- nlsBoot(model)

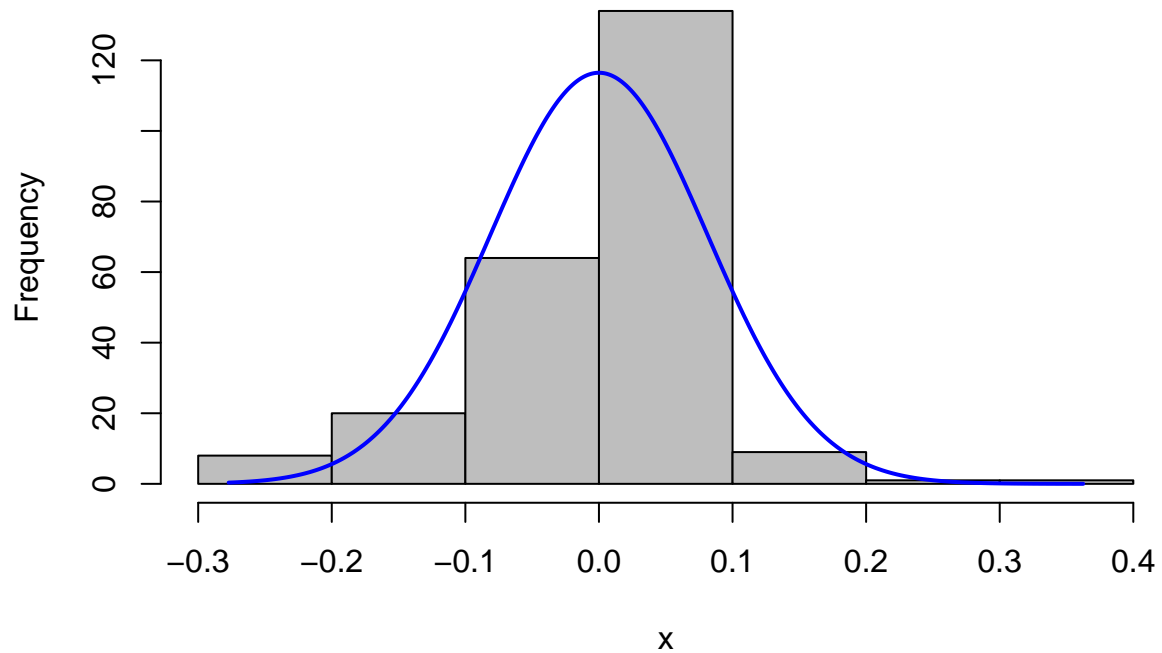
summary(Boot)

##
## -----
## Bootstrap statistics
##           Estimate Std. error
## a      0.21842721 0.037845938
## b      0.01323534 0.001087453
## clx 112.72543911 4.725630630
##
## -----
## Median of bootstrap estimates and percentile confidence intervals
##           Median           2.5%           97.5%
## a      0.2195034    0.13939037    0.28764331
## b      0.0131886    0.01126011    0.01548387
## clx 112.7423863 103.67475871 122.03782922

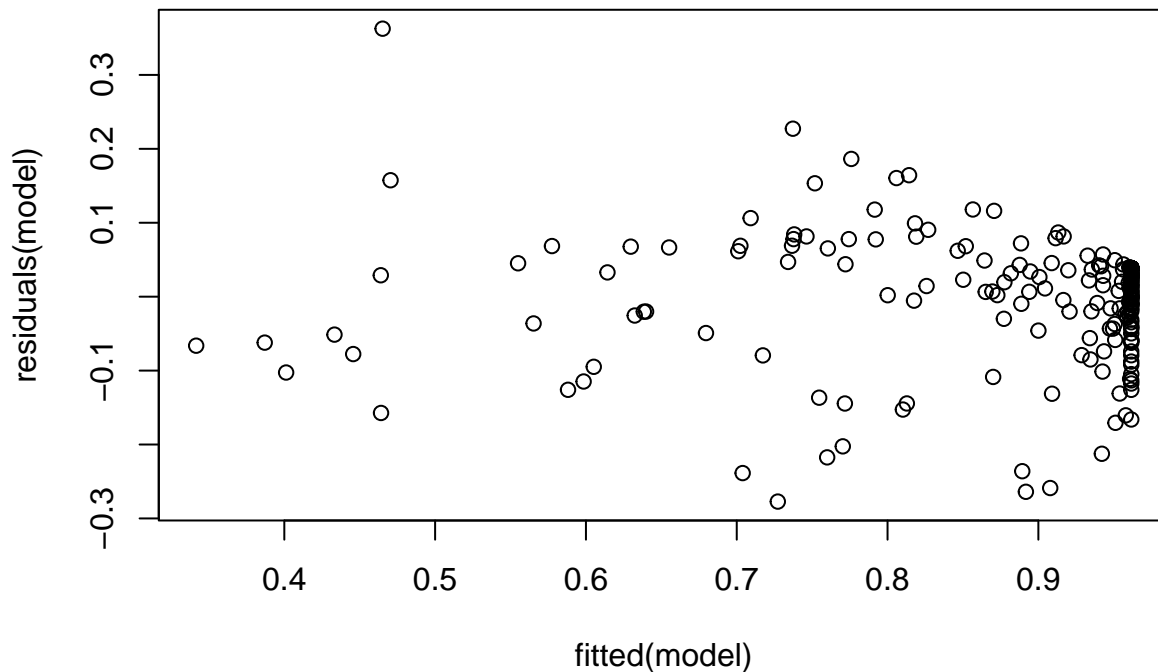
plotPredy(data = paper3_gsdata,
           x     = PI_N_Uptake,
           y     = SI,
           model = model,
           xlab  = "PI Total N Uptake",
           ylab  = "GreenSeeker NDVI SI")
```



```
x <- residuals(model)
plotNormalHistogram(x)
```



```
plot(fitted(model),
     residuals(model))
```



```
a <- summary(model)$coefficients[1]
b <- summary(model)$coefficients[2]
clx <- summary(model)$coefficients[3]
```

```
plateau <- a + b * clx + (-0.5*b)*clx
plateau
```

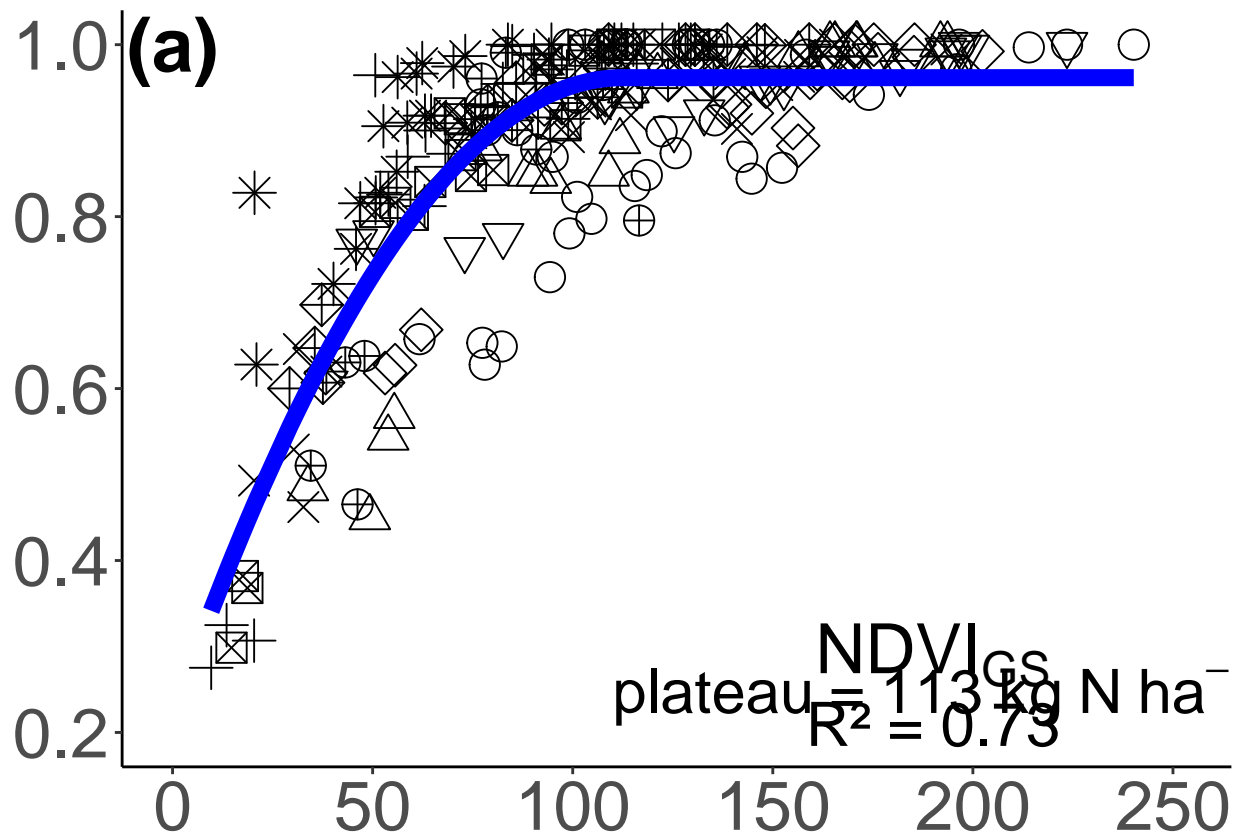
```
## [1] 0.9616152
```

```
GS_ndvi_fit_quadplat <- fitted(model)
```

```
GS_ndvi_quadplat_df <- data.frame(paper3_gsdata$PI_N_Uptake , GS_ndvi_fit_quadplat) #creates dataframe
```

```
GS_ndvi_si_quad_plot <- ggplot(data = paper3_gsdata , aes ( x = PI_N_Uptake , y = SI)) +
  geom_point(data = paper3_gsdata , aes ( x = PI_N_Uptake , y = SI , shape = site_year) , size = 5) +
  geom_line(data = GS_ndvi_quadplat_df , aes( x = paper3_gsdata.PI_N_Uptake , y = GS_ndvi_fit_quadplat)) +
  coord_cartesian(ylim = c(0.20 , 1) , xlim = c(0 , 252)) +
  scale_x_continuous(breaks = seq(0 , 250 , by = 50)) +
  theme_classic() +
  labs(x = NULL , y = NULL) +
  annotate("text", x = 0 , y = 1, label = "(a)", size = 10 , color = "black" , hjust = .5 , fontface = "bold") +
  annotate("text", x = 190 , y = .29, label = "NDVI[GS]", size = 9 , color = "black" , hjust = .5 , fontface = "bold") +
  annotate("text", x = 190 , y = .25, label = "plateau~='~113~kg~N~ha~-1'", size = 8 , color = "black" , hjust = .5 , fontface = "bold") +
  annotate("text", x = 190 , y = .21, label = "R² = 0.73", size = 8 , color = "black" , hjust = .5 , fontface = "bold") +
  theme(axis.title = element_text(size = 26),
        axis.text = element_text(size = 26),
        legend.text = element_text(size = 24),
        legend.title = element_text(size = 24)) +
  geom_hline(yintercept = 0 , size = 0.75) +
  theme(legend.position = "none") +
  scale_shape_manual(values = c(1:20))
```

```
GS_ndvi_si_quad_plot
```



suas-ndre-SI

```
fit.lm2 <- lm(SI ~ PI_N_Uptake, data= paper3_uas_ndre_data)

a.ini2 <- fit.lm2$coefficients[1]
b.ini2 <- fit.lm2$coefficients[2]
clx.ini2 <- mean(paper3_uas_ndre_data$PI_N_Uptake)

quadplat <- function(x, a, b, clx) {
  ifelse(x < clx, a + b * x + (-0.5*b/clx) * x * x,
         a + b * clx + (-0.5*b/clx) * clx * clx)}

model2 <- nls(SI ~ quadplat(PI_N_Uptake, a, b, clx),
              data = paper3_uas_ndre_data,
              start = list(a = a.ini2,
                           b = b.ini2,
                           clx = clx.ini2),
              trace = FALSE,
              nls.control(maxiter = 1000))

summary(model2)

##
## Formula: SI ~ quadplat(PI_N_Uptake, a, b, clx)
##
## Parameters:
##      Estimate Std. Error t value Pr(>|t|)
```

```

## a    3.996e-01  2.280e-02  17.52   <2e-16 ***
## b    8.840e-03  5.969e-04  14.81   <2e-16 ***
## clx  1.287e+02  4.863e+00  26.48   <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.05606 on 234 degrees of freedom
##
## Number of iterations to convergence: 7
## Achieved convergence tolerance: 5.347e-06

nullfunct <- function(x, m){m}

m.ini2    <- mean(paper3_uas_ndre_data$SI)

null2 <- nls(SI ~ nullfunct(PI_N_Uptake, m),
             data = paper3_uas_ndre_data,
             start = list(m = m.ini2),
             trace = FALSE,
             nls.control(maxiter = 1000))

nagelkerke(model2,
            null2)

## $Models
##
## Model: "nls, SI ~ quadplat(PI_N_Uptake, a, b, clx), paper3_uas_ndre_data, list(a = a.ini2, b = b.ini2)"
## Null:  "nls, SI ~ nullfunct(PI_N_Uptake, m), paper3_uas_ndre_data, list(m = m.ini2), list(1000, 1e-06)"
##
## $Pseudo.R.squared.for.model.vs.null
##                                Pseudo.R.squared
## McFadden                      -1.197260
## Cox and Snell (ML)             0.798212
## Nagelkerke (Cragg and Uhler)   -0.284368
##
## $Likelihood.ratio.test
## Df.diff LogLik.diff Chisq    p.value
##      -2      -189.66 379.33 4.2675e-83
##
## $Number.of.observations
##
## Model: 237
## Null:  237
##
## $Messages
## [1] "Note: For models fit with REML, these statistics are based on refitting with ML"
##
## $Warnings
## [1] "None"

confint2(model2,
          level = 0.95)

##           2.5 %           97.5 %
## a    0.35471353  0.44456994
## b    0.00766395  0.01001592

```

```
## clx 119.16598951 138.32655129
```

```
Boot2 <- nlsBoot(model2)
```

```
summary(Boot2)
```

```
##
```

```
## -----
```

```
## Bootstrap statistics
```

```
##      Estimate   Std. error
```

```
## a      0.39834727 0.0228277056
```

```
## b      0.00888344 0.0005930768
```

```
## clx 128.67463380 4.8279270730
```

```
##
```

```
## -----
```

```
## Median of bootstrap estimates and percentile confidence intervals
```

```
##      Median      2.5%      97.5%
```

```
## a  3.998598e-01 3.511902e-01  0.44259004
```

```
## b  8.855489e-03 7.793527e-03  0.01009698
```

```
## clx 1.284738e+02 1.199725e+02 138.73898403
```

```
plotPredy(data = paper3_uas_ndre_data,
```

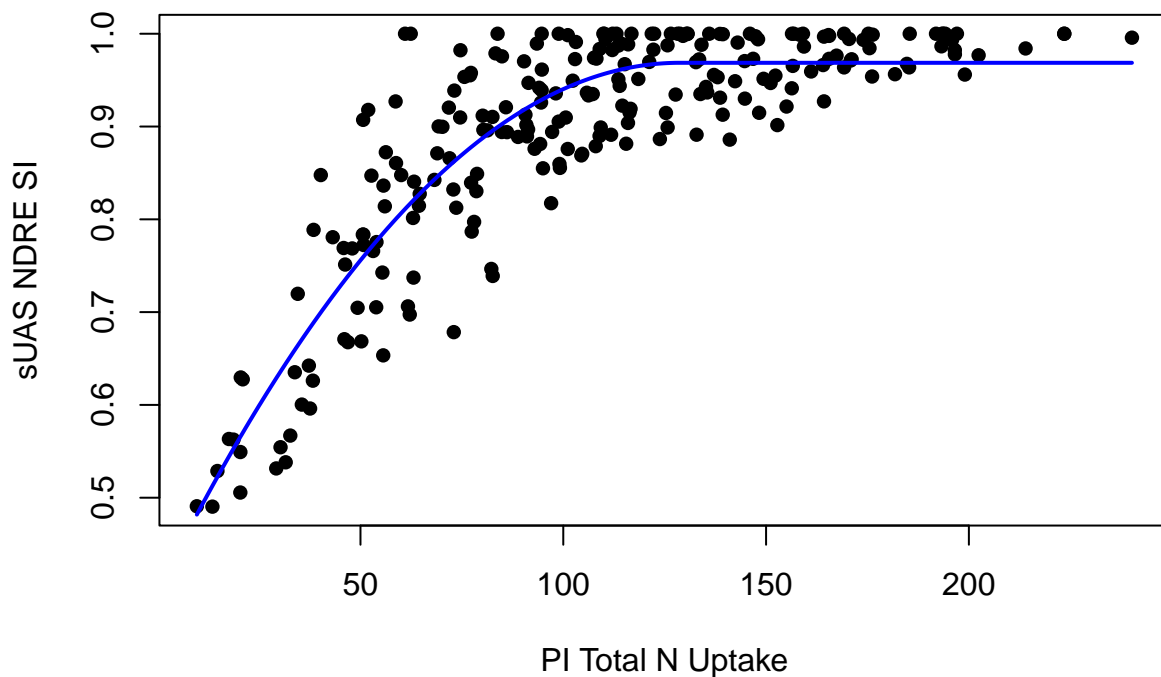
```
  x = PI_N_Uptake,
```

```
  y = SI,
```

```
  model = model2,
```

```
  xlab = "PI Total N Uptake",
```

```
  ylab = "sUAS NDRE SI")
```



```
a2 <- summary(model2)$coefficients[1]
```

```
b2 <- summary(model2)$coefficients[2]
```

```
clx2 <- summary(model2)$coefficients[3]
```

```
plateau2 <- a2 + b2 * clx2 + (-0.5*b2)*clx2
```

```
plateau2
```

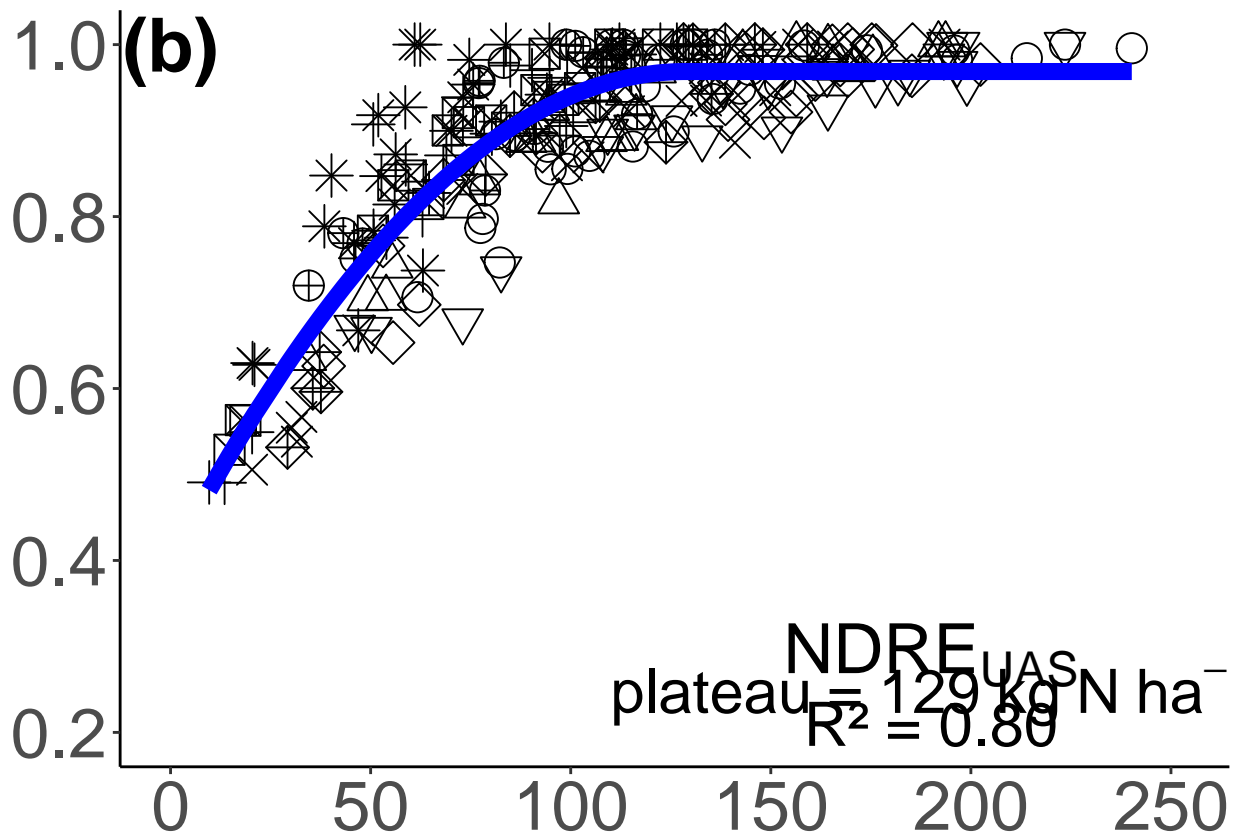
```
## [1] 0.9686962
```

```
sUAS_ndre_fit_quadplat <- fitted(model2)
```

```
sUAS_ndre_quadplat_df <- data.frame(paper3_uas_ndre_data$PI_N_Uptake , sUAS_ndre_fit_quadplat) #creates
```

```
suas_ndre_si_quad_plot <- ggplot(data = paper3_uas_ndre_data , aes ( x = PI_N_Uptake , y = SI)) +
  geom_point(data = paper3_uas_ndre_data , aes ( x = PI_N_Uptake , y = SI , shape = site_year) , size =
  geom_line(data = sUAS_ndre_quadplat_df , aes( x = paper3_uas_ndre_data.PI_N_Uptake , y = sUAS_ndre_fi
  coord_cartesian(ylim = c(0.20 , 1) , xlim = c(0 , 252)) +
  scale_x_continuous(breaks = seq(0 , 250 , by = 50)) +
  theme_classic() +
  labs(x = NULL , y = NULL) +
  annotate("text", x = 0 , y = 1, label = "(b)", size = 10 , color = "black" , hjust = 0.5 , fontface =
  annotate("text", x = 190 , y = .29, label = "NDRE[UAS]", size = 9 , color = "black" , hjust = .5 , pa
  annotate("text", x = 190 , y = .25, label = "plateau~'~129~kg~N~ha~-1", size = 8 , color = "black"
  annotate("text", x = 190 , y = .21, label = "R² = 0.80", size = 8 , color = "black" , hjust = .5) +
  theme(axis.title = element_text(size = 26),
        axis.text = element_text(size = 26),
        legend.text = element_text(size = 24),
        legend.title = element_text(size = 24)
  ) +
  geom_hline(yintercept = 0 , size = 0.75) +
  theme(legend.position = "none") +
  scale_shape_manual(values = c(1:20))
```

```
suas_ndre_si_quad_plot
```



suas-ndvi-SI

```
fit.lm3 <- lm(SI ~ PI_N_Uptake, data= paper3_uas_ndvi_data)
```

```
a.ini3 <- fit.lm3$coefficients[1]
```

```
b.ini3 <- fit.lm3$coefficients[2]
```

```
clx.ini3 <- mean(paper3_uas_ndvi_data$PI_N_Uptake)
```

```
quadplat <- function(x, a, b, clx) {
  ifelse(x < clx, a + b * x + (-0.5*b/clx) * x * x,
         a + b * clx + (-0.5*b/clx) * clx * clx)}
```

```
model3 <- nls(SI ~ quadplat(PI_N_Uptake, a, b, clx),
  data = paper3_uas_ndvi_data,
  start = list(a = a.ini3,
               b = b.ini3,
               clx = clx.ini3),
  trace = FALSE,
  nls.control(maxiter = 1000))
```

```
summary(model3)
```

```
##
```

```
## Formula: SI ~ quadplat(PI_N_Uptake, a, b, clx)
```

```
##
```

```
## Parameters:
```

```
##      Estimate Std. Error t value Pr(>|t|)
```

```
## a    6.458e-01  1.403e-02   46.02  <2e-16 ***
```

```
## b    7.078e-03  4.723e-04   14.99  <2e-16 ***
```

```
## clx  9.646e+01  3.180e+00   30.34  <2e-16 ***
```

```
## ---
```

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

```
##
```

```
## Residual standard error: 0.02632 on 234 degrees of freedom
```

```
##
```

```
## Number of iterations to convergence: 11
```

```
## Achieved convergence tolerance: 7.936e-06
```

```
nullfunct <- function(x, m){m}
```

```
m.ini3 <- mean(paper3_uas_ndvi_data$SI)
```

```
null3 <- nls(SI ~ nullfunct(PI_N_Uptake, m),
  data = paper3_uas_ndvi_data,
  start = list(m = m.ini3),
  trace = FALSE,
  nls.control(maxiter = 1000))
```

```
nagelkerke(model3,
  null3)
```

```
## $Models
```

```
##
```

```
## Model: "nls, SI ~ quadplat(PI_N_Uptake, a, b, clx), paper3_uas_ndvi_data, list(a = a.ini3, b = b.ini3)"
```

```
## Null: "nls, SI ~ nullfunct(PI_N_Uptake, m), paper3_uas_ndvi_data, list(m = m.ini3), list(1000, 1e-06)"
```



```
##
## $Pseudo.R.squared.for.model.vs.null
##                                Pseudo.R.squared
## McFadden                      -0.605012
## Cox and Snell (ML)             0.813116
## Nagelkerke (Cragg and Uhler)   -0.054225
##
## $Likelihood.ratio.test
## Df.diff LogLik.diff Chisq    p.value
##      -2      -198.76 397.51 4.7998e-87
##
## $Number.of.observations
##
## Model: 237
## Null: 237
##
## $Messages
## [1] "Note: For models fit with REML, these statistics are based on refitting with ML"
##
## $Warnings
## [1] "None"

confint2(model3,
          level = 0.95)

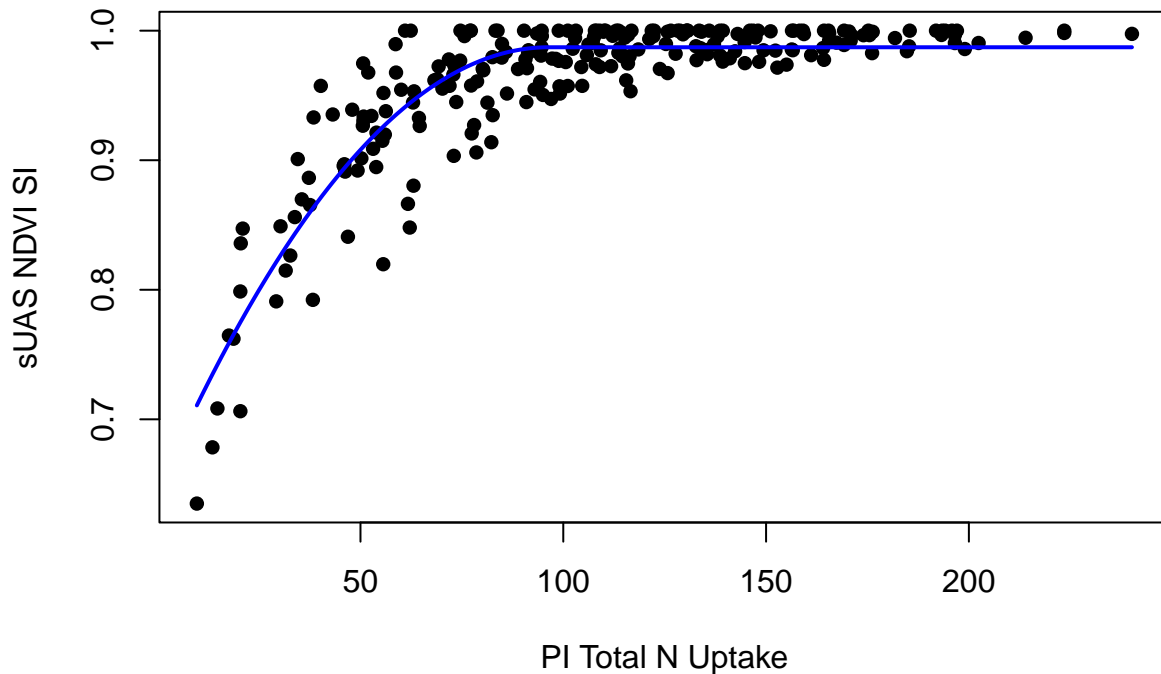
##           2.5 %           97.5 %
## a    0.618157809 6.734574e-01
## b    0.006147996 8.008987e-03
## clx 90.196205891 1.027258e+02

Boot3 <- nlsBoot(model3)

summary(Boot3)

##
## -----
## Bootstrap statistics
##      Estimate Std. error
## a    0.64458101 0.0140131539
## b    0.00712298 0.0004710038
## clx 96.40076649 3.1039897943
##
## -----
## Median of bootstrap estimates and percentile confidence intervals
##           Median           2.5%           97.5%
## a    0.645410303 0.616728996 6.698013e-01
## b    0.007100738 0.006275563 8.070276e-03
## clx 96.344508975 90.543489243 1.026824e+02

plotPredy(data = paper3_uas_ndvi_data,
           x    = PI_N_Uptake,
           y    = SI,
           model = model3,
           xlab = "PI Total N Uptake",
           ylab = "sUAS NDVI SI")
```



```
a3 <- summary(model3)$coefficients[1]
b3 <- summary(model3)$coefficients[2]
clx3 <- summary(model3)$coefficients[3]

plateau3 <- a3 + b3 * clx3 + (-0.5*b3)*clx3
plateau3
```

```
## [1] 0.9872068
```

```
sUAS_ndvi_fit_quadplat <- fitted(model3)
```

```
sUAS_ndvi_quadplat_df <- data.frame(paper3_uas_ndvi_data$PI_N_Uptake , sUAS_ndvi_fit_quadplat) #creates
```

```
suas_ndvi_si_quad_plot <- ggplot(data = paper3_uas_ndvi_data , aes ( x = PI_N_Uptake , y = SI)) +
  geom_point(data = paper3_uas_ndvi_data , aes ( x = PI_N_Uptake , y = SI , shape = site_year) , size =
  geom_line(data = sUAS_ndvi_quadplat_df , aes( x = paper3_uas_ndvi_data.PI_N_Uptake , y = sUAS_ndvi_fit_quadplat)) +
  coord_cartesian(ylim = c(0.20 , 1) , xlim = c(0 , 252)) +
  scale_x_continuous(breaks = seq(0 , 250 , by = 50)) +
  theme_classic() +
  labs(x = NULL , y = NULL , shape = "Site-Year") +
  annotate("text", x = 0 , y = 1, label = "(c)", size = 10 , color = "black" , hjust = 0.5 , fontface = "bold") +
  annotate("text", x = 190 , y = .29, label = "NDVI[UAS]", size = 9 , color = "black" , hjust = .5 , fontface = "bold") +
  annotate("text", x = 190 , y = .25, label = "plateau~='~96~kg~N~ha^-1'", size = 8 , color = "black" , hjust = .5 , fontface = "bold") +
  annotate("text", x = 190 , y = .21, label = "R² = 0.81", size = 8 , color = "black" , hjust = .5 , fontface = "bold") +
  theme(axis.title = element_text(size = 26),
        axis.text = element_text(size = 26),
        legend.text = element_text(size = 20),
        legend.title = element_text(size = 20 , hjust = 0.5),
        legend.box.background = element_rect(size = 1)) +
  geom_hline(yintercept = 0 , size = 0.75) +
  theme(legend.position = c(.72 , .4)) +
  scale_shape_manual(values = c(1:20))
```

```
suas_ndvi_si_quad_plot
```

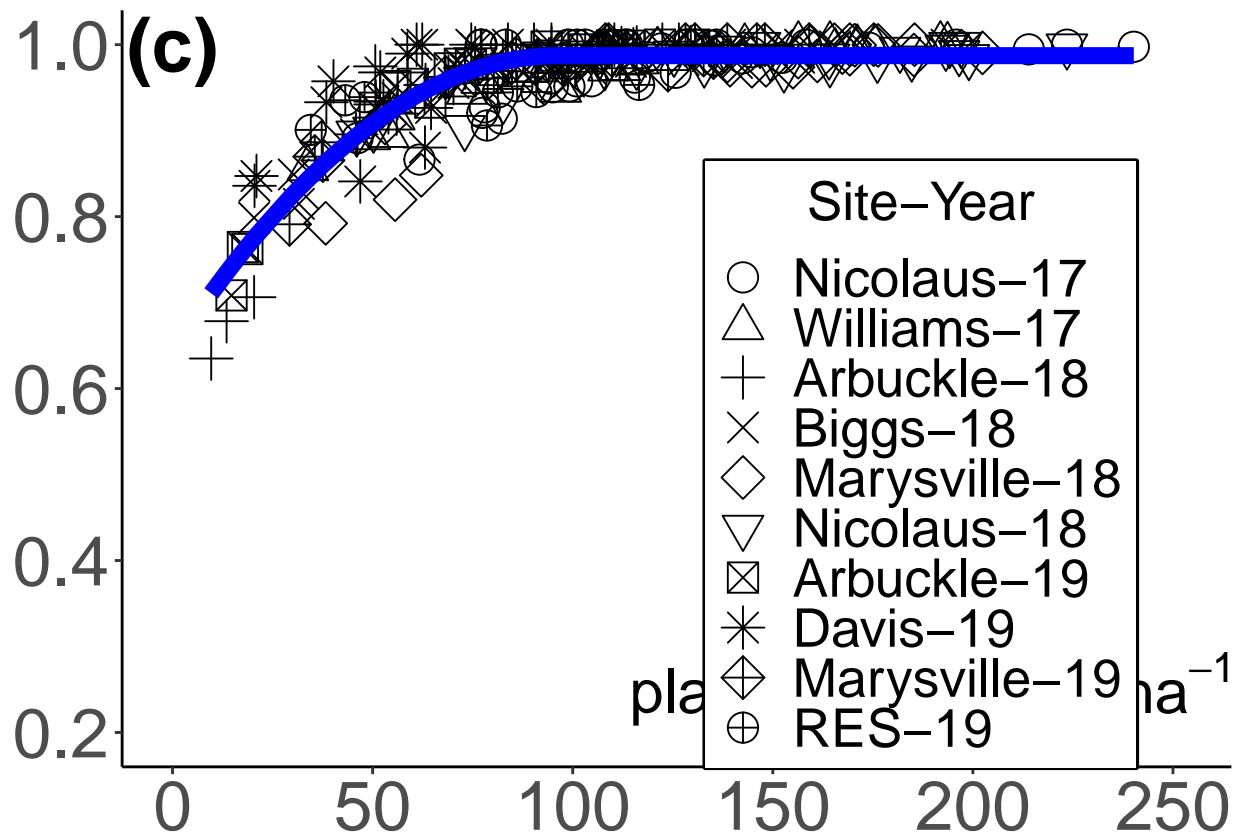
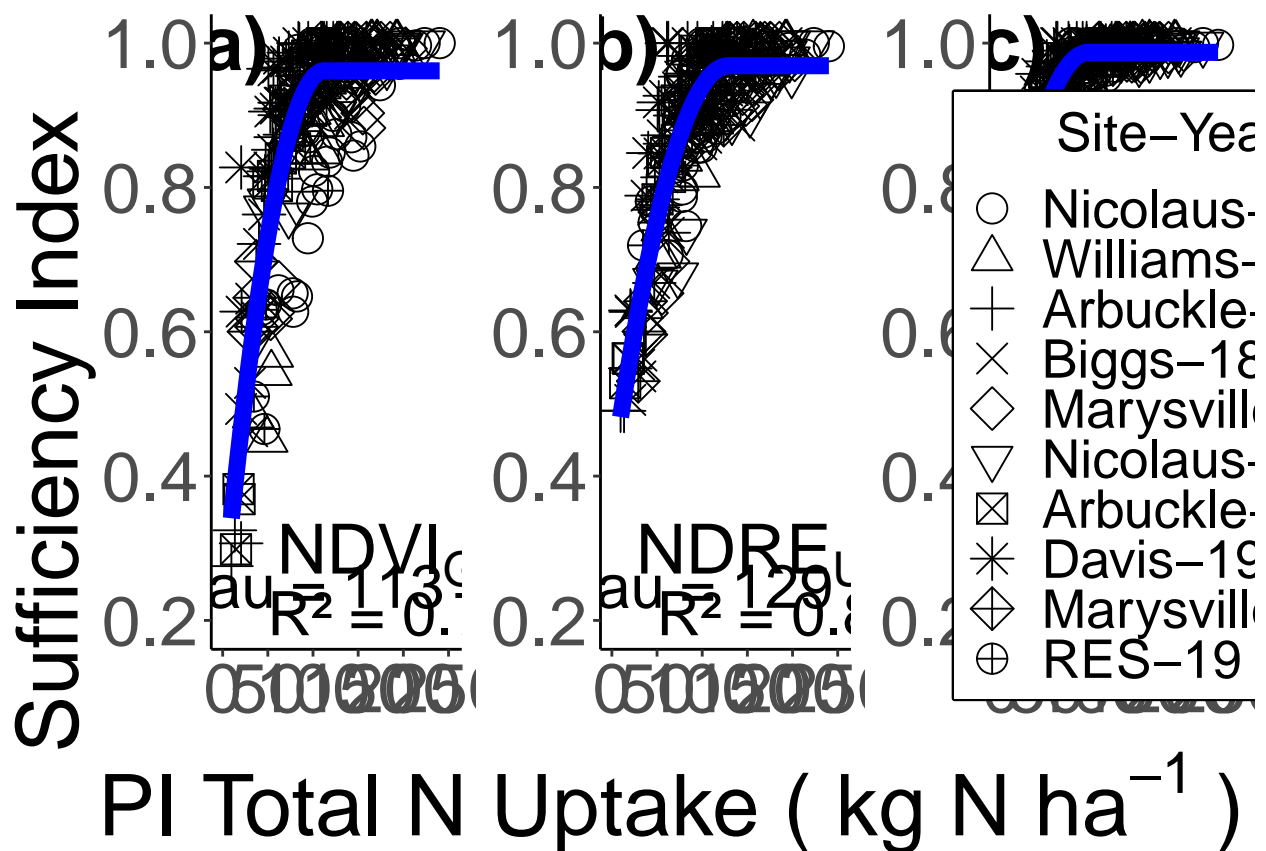


FIGURE 4

```
Fig4 <- grid.arrange(arrangeGrob(GS_ndvi_si_quad_plot,
                                suas_ndre_si_quad_plot,
                                suas_ndvi_si_quad_plot,
                                left = textGrob("Sufficiency Index" , rot = 90 , gp = gpar(fontsize = 12)),
                                bottom = textGrob("PI Total N Uptake ( kg N ha-1~")" , gp = gpar(fontsize = 12)),
                                nrow = 1,
                                ncol = 3
                                ))
```



```
ggsave("FIGURES/Fig4.tiff" , Fig4 , compression = "lzw" , width = 22 , height = 10, type = "cairo" , dp
```

FIGURE 5

gs ndvi SI

```
ctrl <- lmeControl(opt = "optim") #changes control to "optimal" settings
```

```
gsndvi_yield_mixlm <- lme(fixed = GrainYield_Mgha ~ SI ,
  random = ~I(SI) | site_year,
  data = paper3_gsdata)
```

```
summary(gsndvi_yield_mixlm)
```

model

```
## Linear mixed-effects model fit by REML
## Data: paper3_gsdata
##      AIC      BIC    logLik
## 632.2034 652.9609 -310.1017
##
## Random effects:
## Formula: ~I(SI) | site_year
## Structure: General positive-definite, Log-Cholesky parametrization
##              StdDev    Corr
```

```

## (Intercept) 2.8528028 (Intr)
## I(SI)      3.2215774 -0.936
## Residual   0.7974361
##
## Fixed effects: GrainYield_Mgha ~ SI
##              Value Std.Error DF  t-value p-value
## (Intercept) 1.756200 0.9680527 226 1.814157  0.071
## SI          9.547127 1.0902077 226 8.757164  0.000
## Correlation:
##   (Intr)
## SI -0.943
##
## Standardized Within-Group Residuals:
##      Min      Q1      Med      Q3      Max
## -3.34438501 -0.54121956  0.01143244  0.58140764  2.20039027
##
## Number of Observations: 237
## Number of Groups: 10
summary(gsndvi_yield_mixlm)$tTable

##              Value Std.Error DF  t-value      p-value
## (Intercept) 1.756200 0.9680527 226 1.814157 7.097962e-02
## SI          9.547127 1.0902077 226 8.757164 4.774230e-16
Anova(gsndvi_yield_mixlm , type = 2)

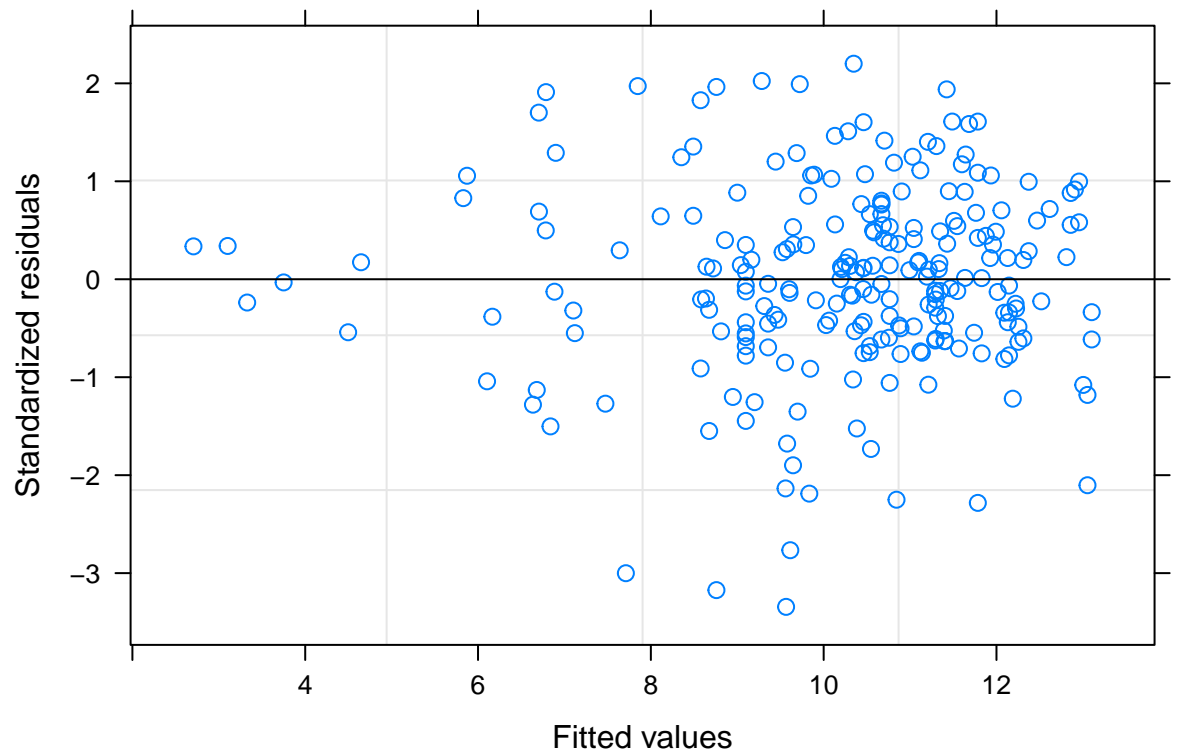
## Analysis of Deviance Table (Type II tests)
##
## Response: GrainYield_Mgha
##      Chisq Df Pr(>Chisq)
## SI 76.688  1  < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Anova(gsndvi_yield_mixlm , type = 3)

## Analysis of Deviance Table (Type III tests)
##
## Response: GrainYield_Mgha
##      Chisq Df Pr(>Chisq)
## (Intercept) 3.2912  1  0.06965 .
## SI          76.6879  1  < 2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
r.squaredGLMM(gsndvi_yield_mixlm)

## Warning: 'r.squaredGLMM' now calculates a revised statistic. See the help page.
##      R2m      R2c
## [1,] 0.5345773 0.8466198

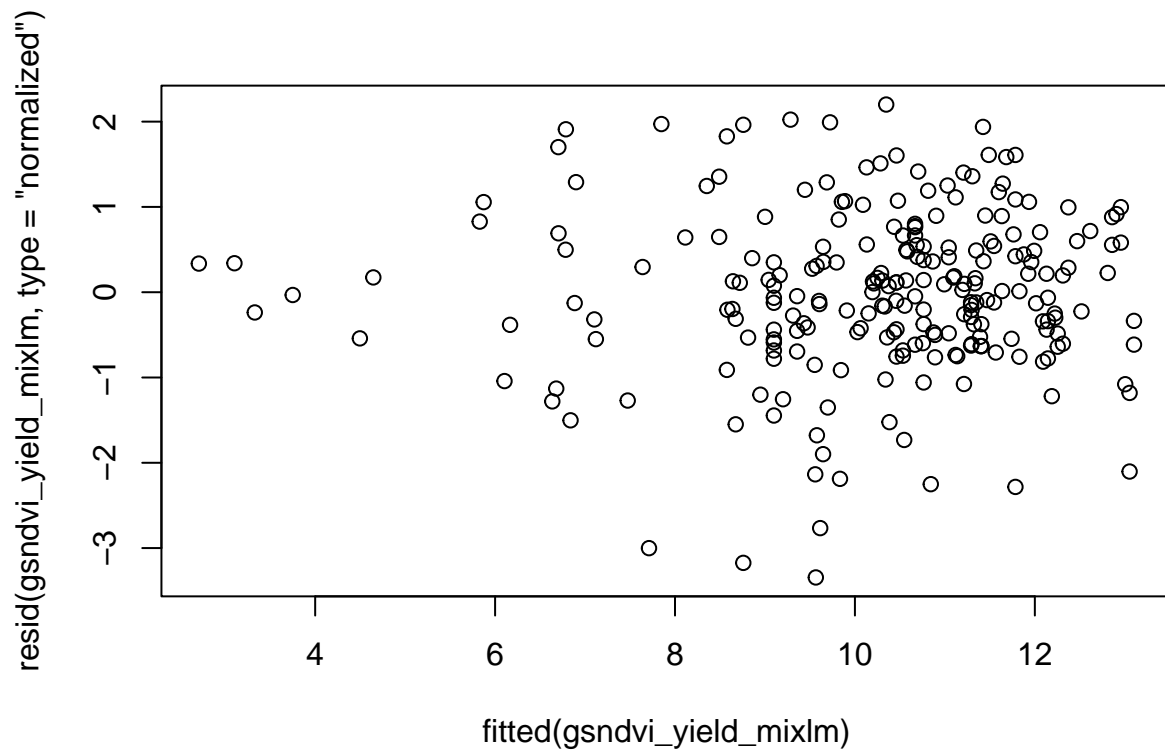
plot (gsndvi_yield_mixlm)

```



diagnostics

```
plot(resid(gsndvi_yield_mixlm, type = "normalized") ~fitted(gsndvi_yield_mixlm))
```

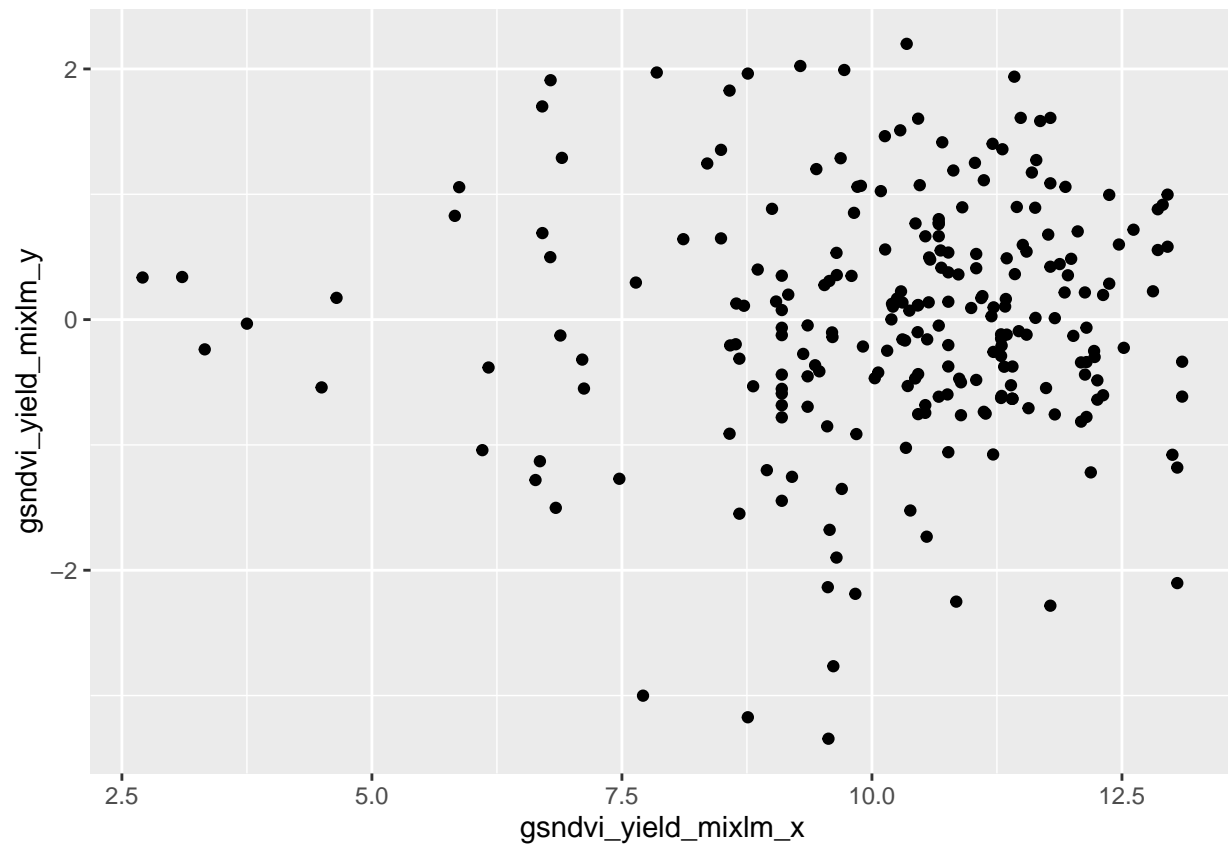


```
gsndvi_yield_mixlm_y <- resid(gsndvi_yield_mixlm, type = "normalized")
```

```
gsndvi_yield_mixlm_x <- fitted(gsndvi_yield_mixlm)
```

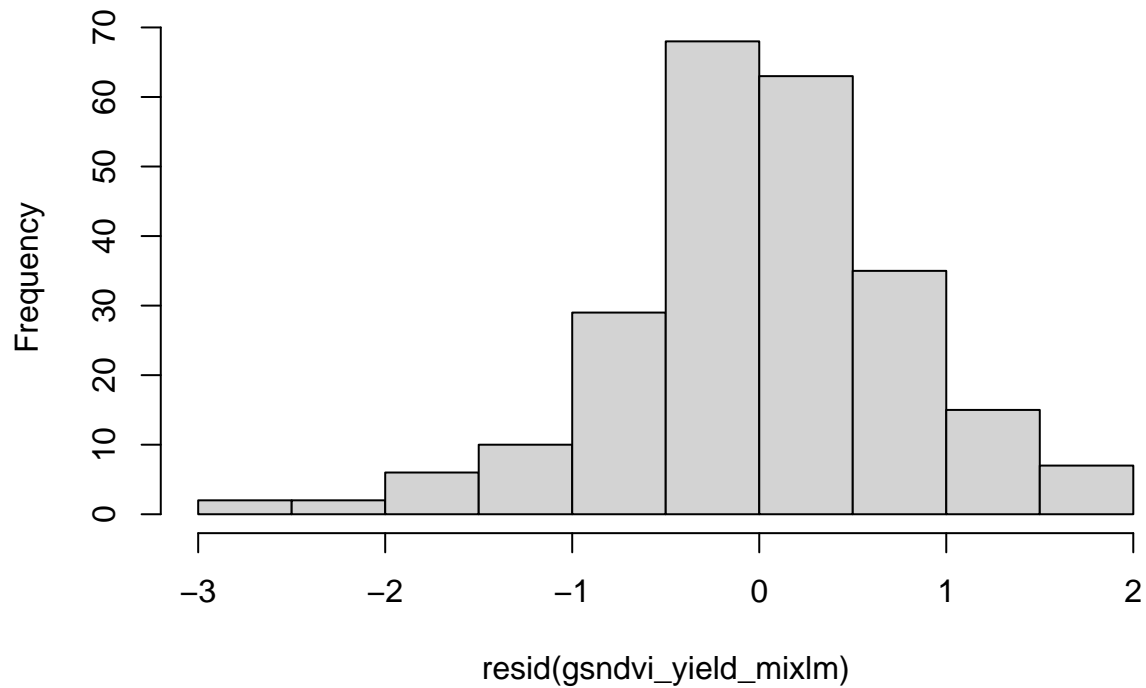
```
gsndvi_yield_mixlmresid_data <- data.frame(gsndvi_yield_mixlm_x , gsndvi_yield_mixlm_y)
```

```
ggplot( data = gsndvi_yield_mixlmresid_data , aes( x = gsndvi_yield_mixlm_x , y = gsndvi_yield_mixlm_y ) )  
  geom_point(mapping = aes( gsndvi_yield_mixlm_x , gsndvi_yield_mixlm_y ) , data = gsndvi_yield_mixlmresid_data )
```

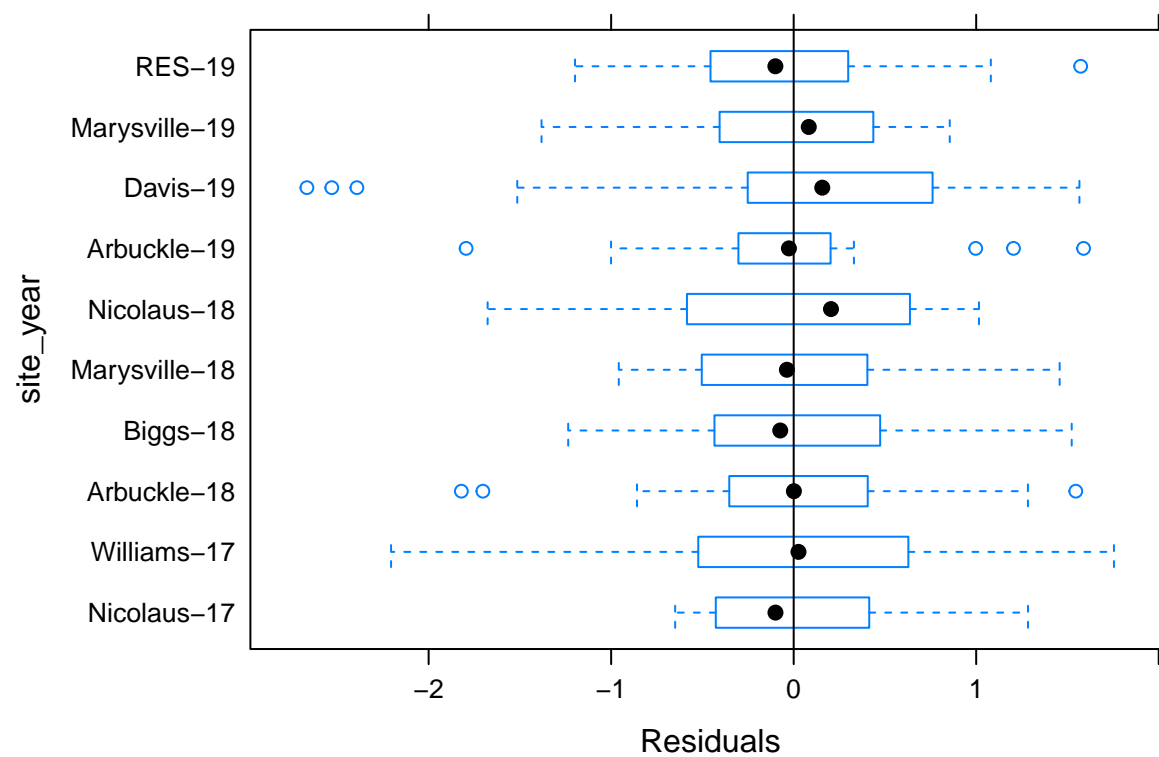


```
hist(resid(gsndvi_yield_mixlm))
```

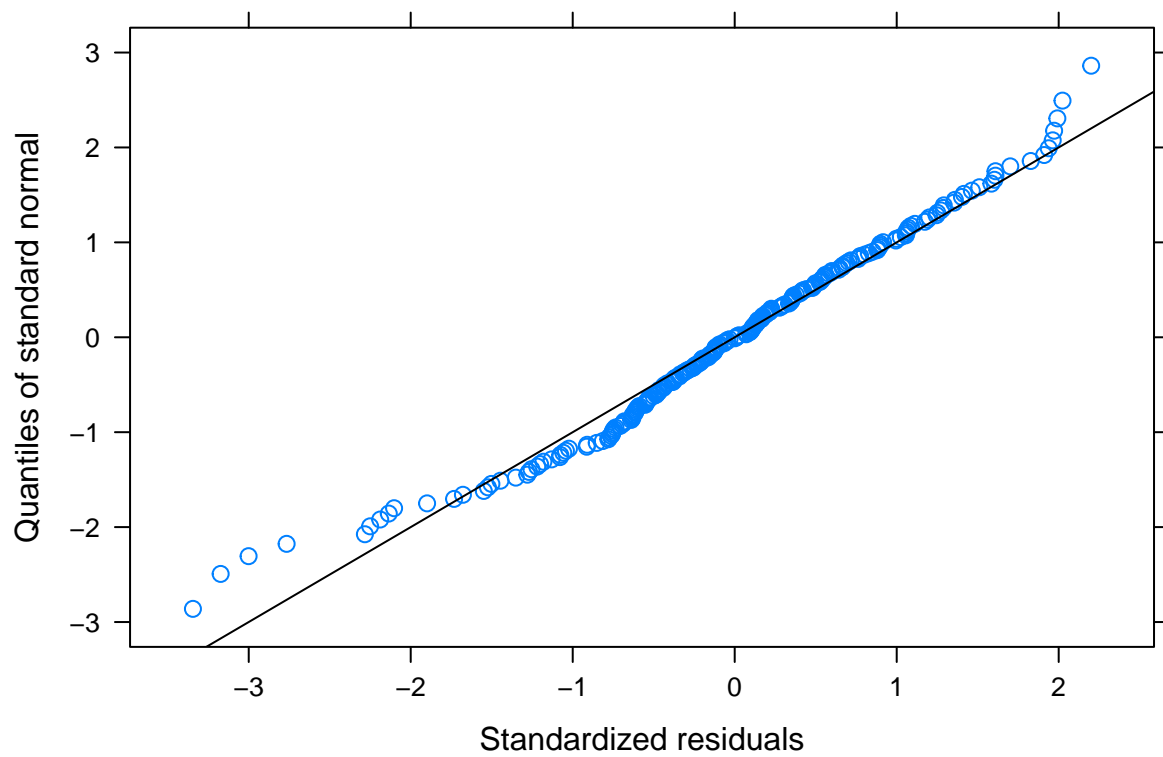
Histogram of resid(gsndvi_yield_mixlm)



```
plot(gsndvi_yield_mixlm, site_year ~ resid(.), abline = 0)
```

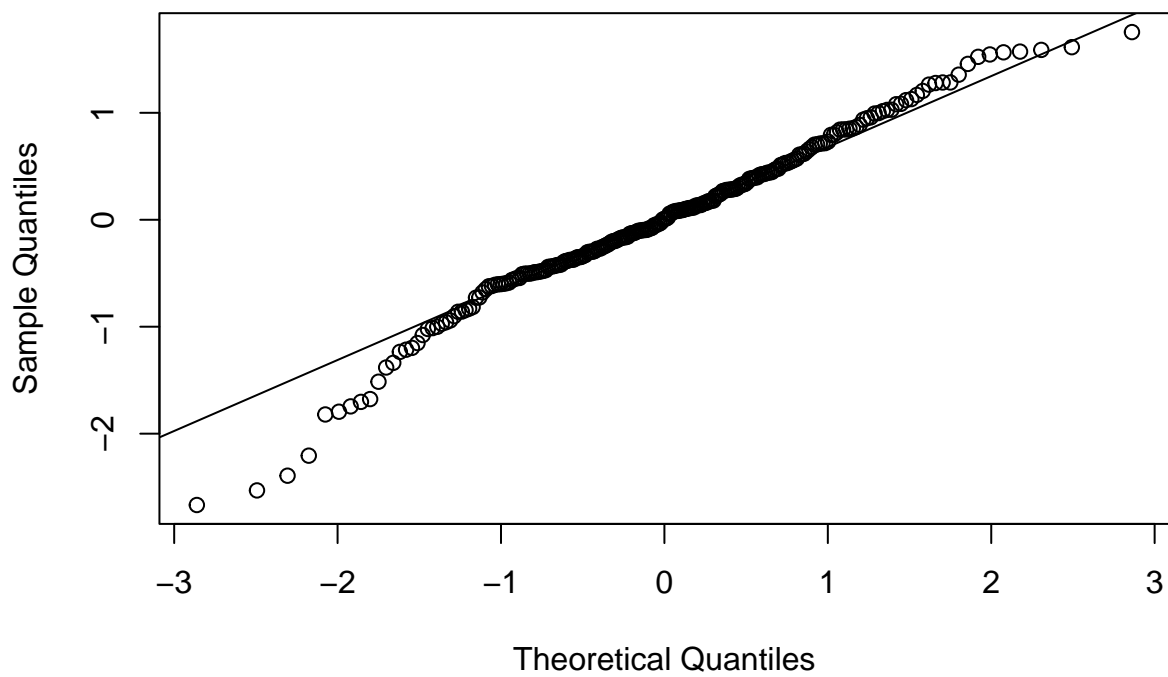


```
qqnorm(gsndvi_yield_mixlm, abline = c(0,1) )
```

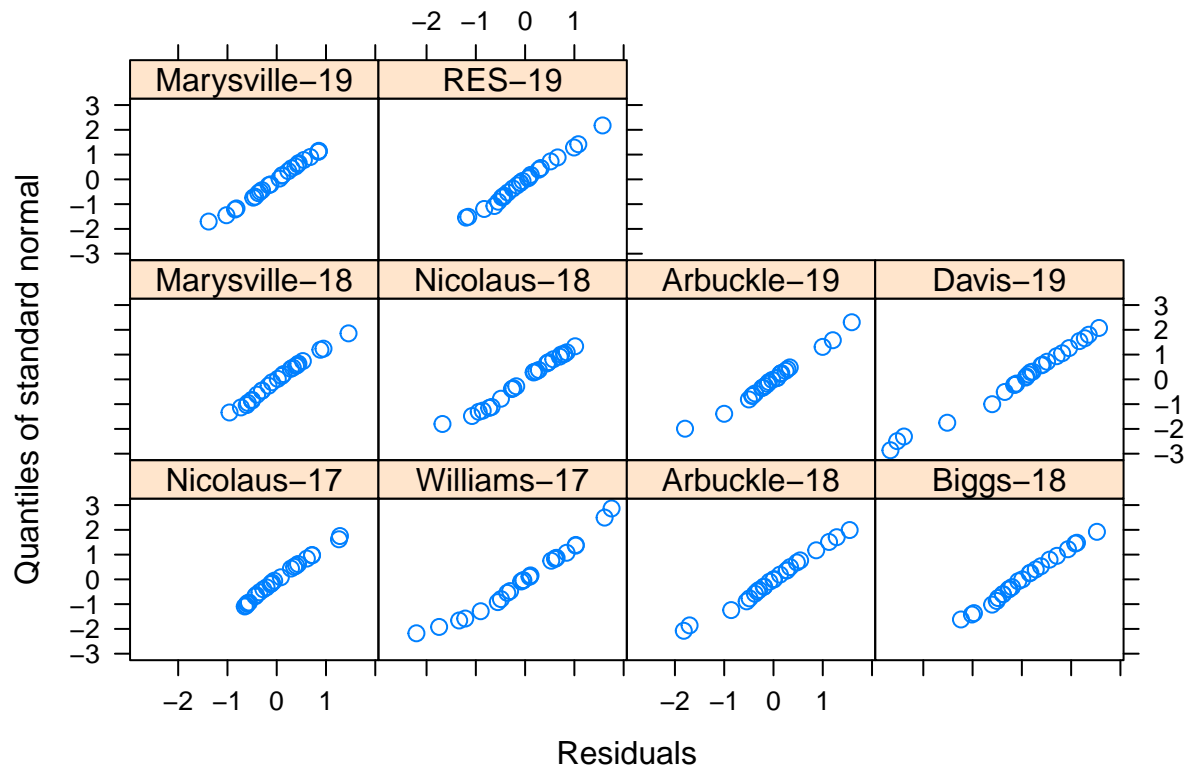



```
qqnorm(resid(gsndvi_yield_mixlm))
qqline(resid(gsndvi_yield_mixlm))
```

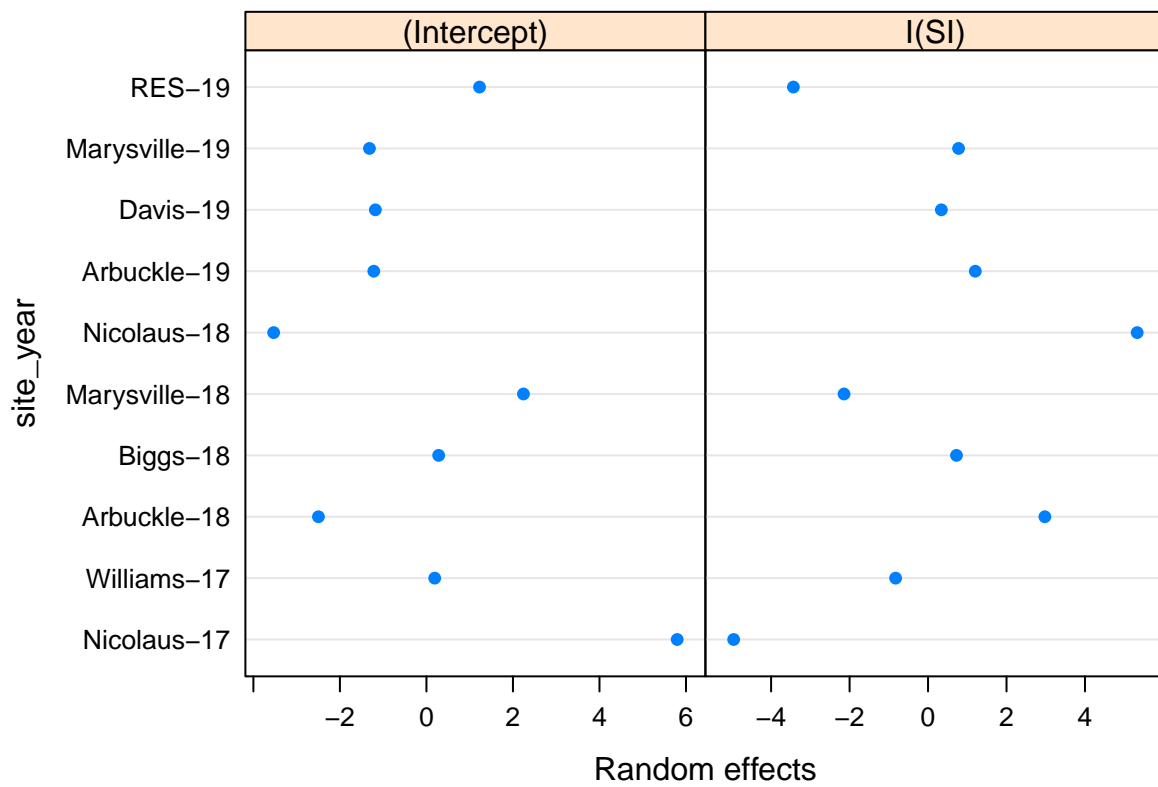
Normal Q–Q Plot



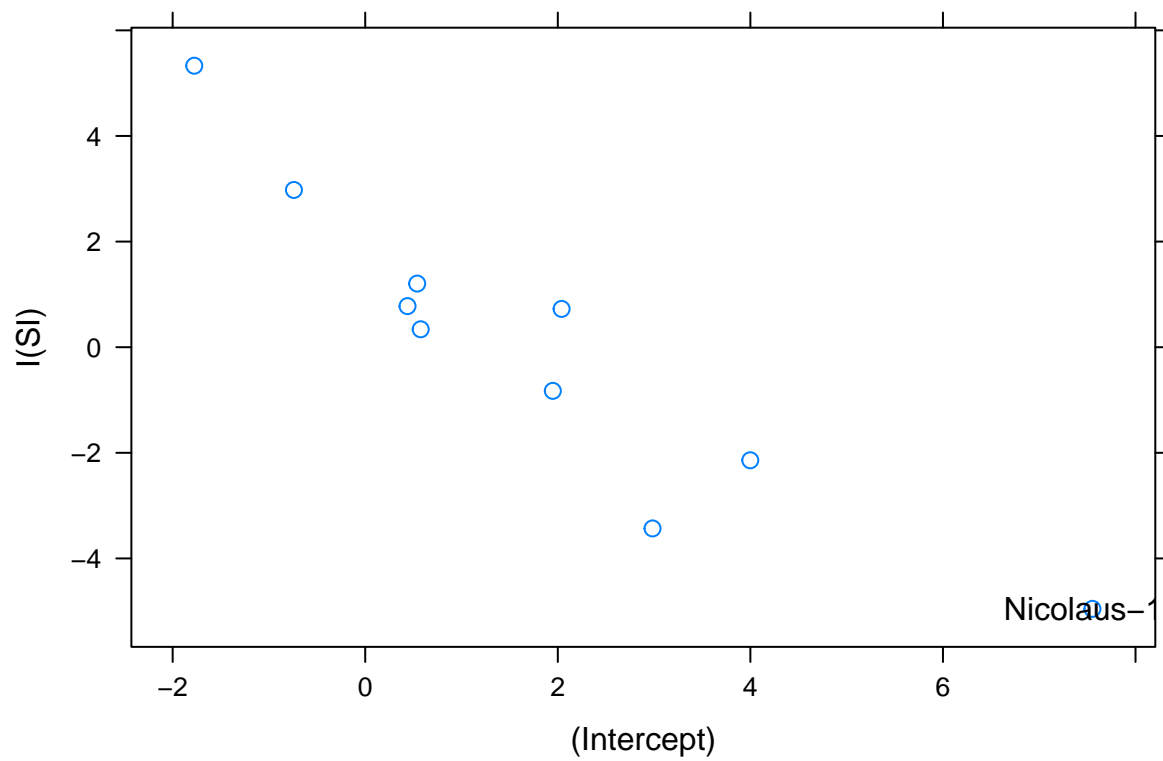
```
qqnorm(gsndvi_yield_mixlm , ~resid(.) | site_year)
```



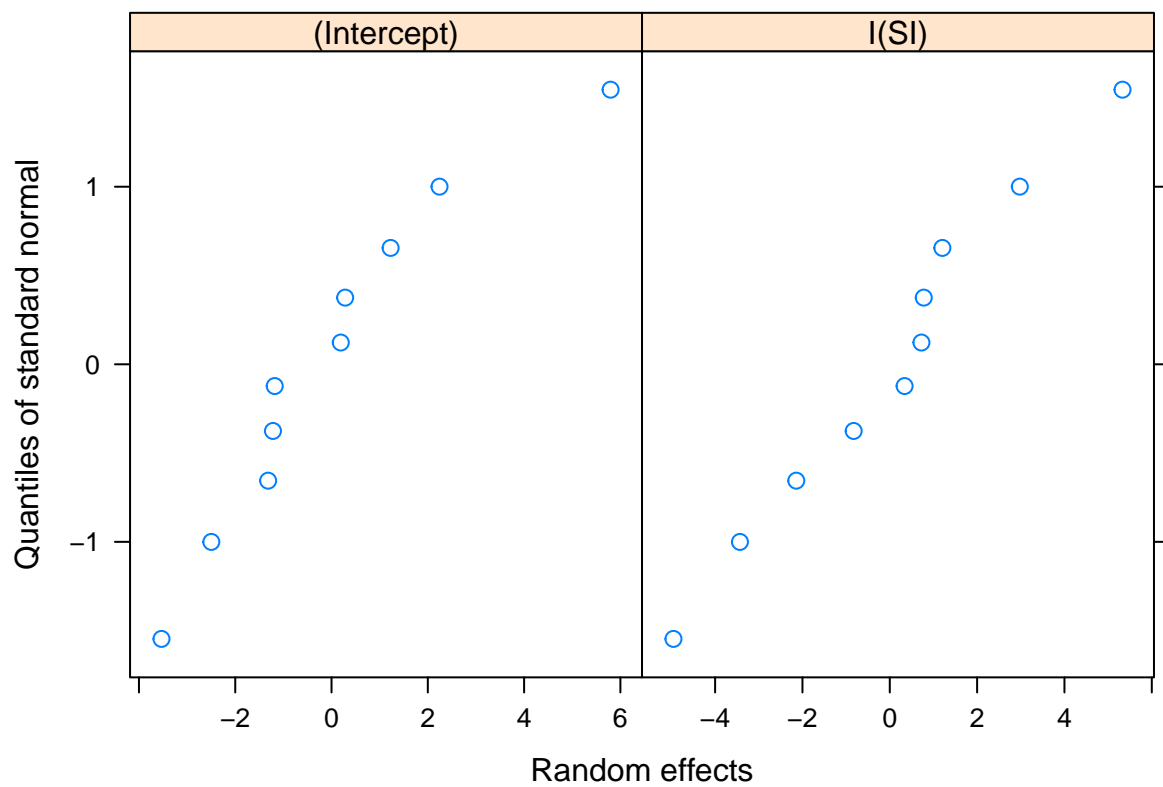
```
plot(ranef(gsndvi_yield_mixlm))
```



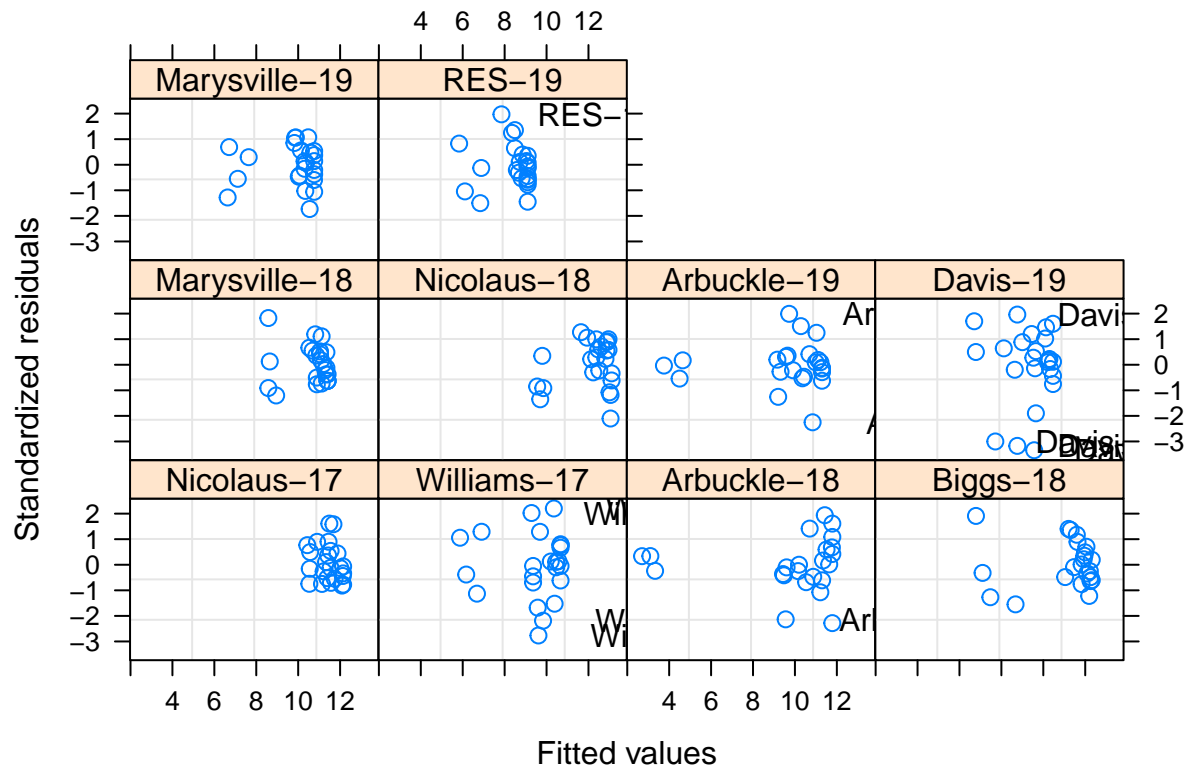
```
pairs(gsndvi_yield_mixlm , id = 0.1)
```



```
qqnorm(gsndvi_yield_mixlm , ~ranef(.))
```



```
plot( gsndvi_yield_mixlm, resid(., type = "p") ~ fitted(.) | site_year,
      id = 0.05, adj = -0.3 )
```



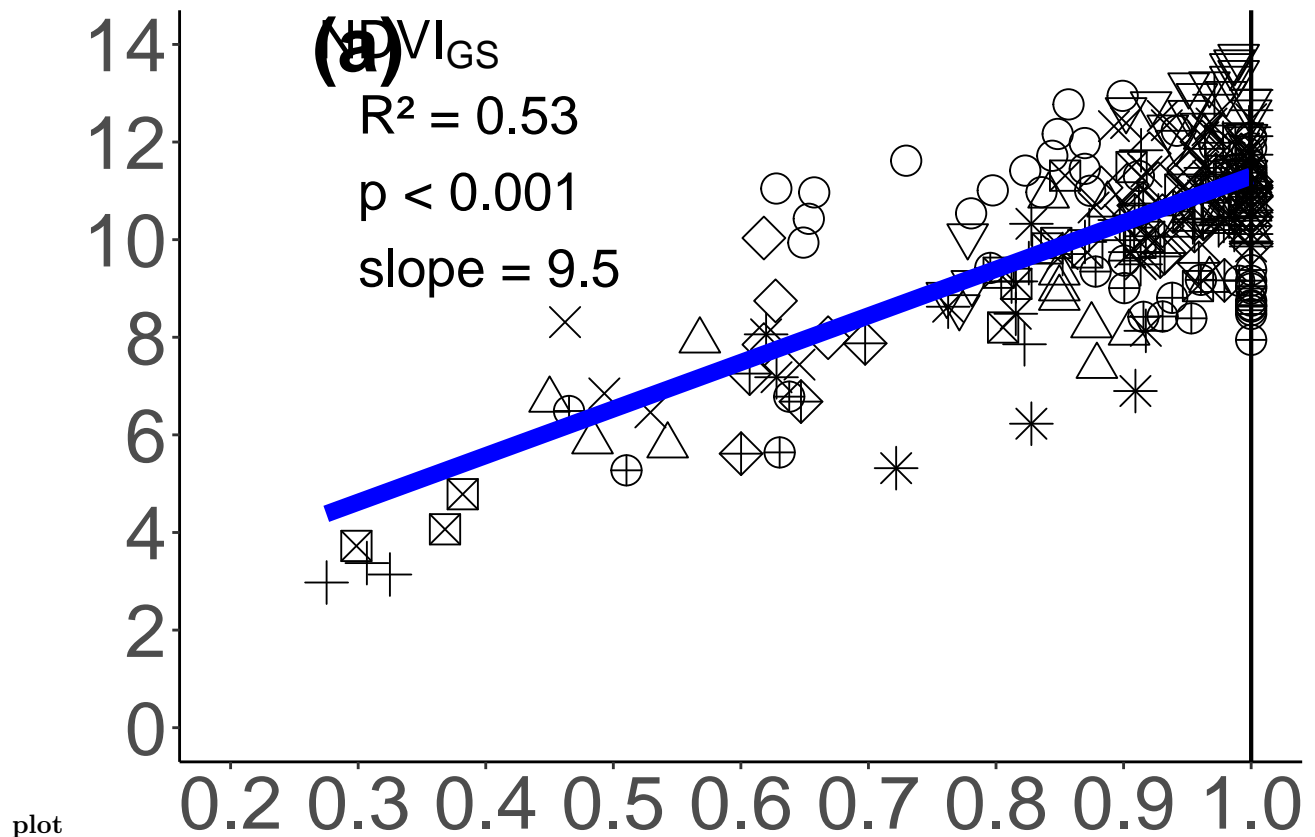
```
mylist <- list(SI=seq(round(min(paper3_gsdata$SI), digits = 3) , 1 , by = .001))

gsndvi_emmeans <- as.data.frame(summary(emmeans(gsndvi_yield_mixlm , ~ SI , at = mylist )))
```

emmeans

```
plot_si1 <- ggplot( data = paper3_gsdata , aes( x = SI , y = GrainYield_Mgha)) +
  geom_point( data = paper3_gsdata , aes( x = SI , y = GrainYield_Mgha , shape = site_year) , size = 5)
  geom_line( data = gsndvi_emmeans , aes(x = SI , y = emmean) , color = "blue" , size = 3) +
  theme_classic() +
  labs( x = NULL , y = NULL , color = "Index") +
  scale_x_continuous(breaks = seq(.2 , 1 , by = .1)) +
  scale_y_continuous(breaks = seq(0 , 14 , by = 2)) +
  coord_cartesian(ylim = c(0,14) , xlim = c(.2 , 1)) +
  geom_vline(xintercept = 1 , size = 0.75) +
  theme(axis.text = element_text(size = 28),
        axis.title = element_text(size = 28),
        legend.text = element_text(size = 24),
        legend.title = element_text(size = 24),
        legend.position = "none") +
  annotate("text", x=0.3, y=11, label="R² = 0.53\np < 0.001\nslope = 9.5", size = 7 , color = "black" ,
  annotate("text", x=0.3, y=14, label="(a)", size = 10 , color = "black" , hjust = 0.5 , fontface = 2)
  annotate("text", x=0.34, y=14, label="NDVI[GS]", size = 7 , color = "black" , hjust = 0.5 , parse = T)
  scale_shape_manual(values = c(1:20))

plot_si1
```



sUAS ndvi RI

```
sUASndvi_yield_mixlm <- lme(fixed = GrainYield_Mgha ~ SI ,
                             random = ~I(SI) | site_year,
                             data = paper3_uas_ndvi_data)
```

```
summary(sUASndvi_yield_mixlm)
```

model

```
## Linear mixed-effects model fit by REML
## Data: paper3_uas_ndvi_data
##      AIC      BIC    logLik
## 639.7817 660.5392 -313.8908
##
## Random effects:
## Formula: ~I(SI) | site_year
## Structure: General positive-definite, Log-Cholesky parametrization
##              StdDev   Corr
## (Intercept) 7.6291818 (Intr)
## I(SI)       8.0549586 -0.99
## Residual    0.8161178
##
## Fixed effects: GrainYield_Mgha ~ SI
##              Value Std.Error DF  t-value p-value
## (Intercept) -14.12294  2.643468 226 -5.342579    0
## SI          25.33529  2.781592 226  9.108198    0
```

```

## Correlation:
## (Intr)
## SI -0.992
##
## Standardized Within-Group Residuals:
##      Min      Q1      Med      Q3      Max
## -5.15225909 -0.55392647  0.05855843  0.56664149  2.24710746
##
## Number of Observations: 237
## Number of Groups: 10
summary(sUASndvi_yield_mixlm)$tTable

##              Value Std.Error DF   t-value      p-value
## (Intercept) -14.12294  2.643468 226 -5.342579 2.232467e-07
## SI           25.33529  2.781592 226  9.108198 4.573975e-17
Anova(sUASndvi_yield_mixlm , type = 2)

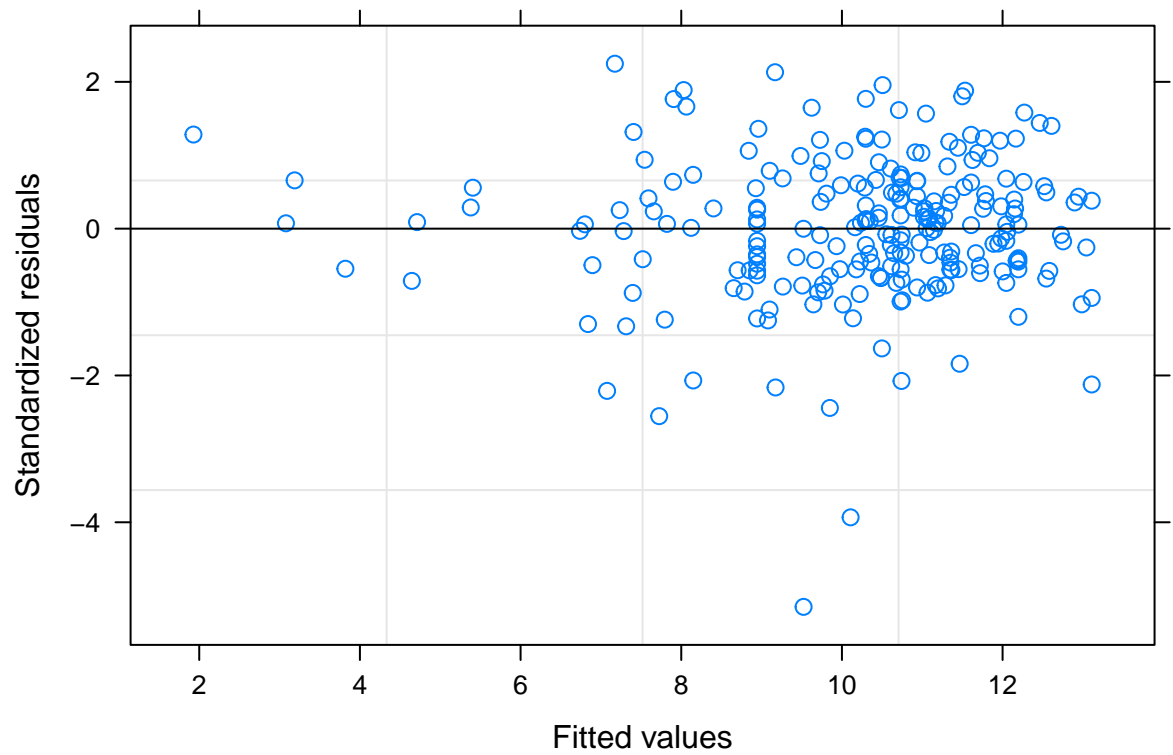
## Analysis of Deviance Table (Type II tests)
##
## Response: GrainYield_Mgha
##      Chisq Df Pr(>Chisq)
## SI 82.959  1  < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Anova(sUASndvi_yield_mixlm , type = 3)

## Analysis of Deviance Table (Type III tests)
##
## Response: GrainYield_Mgha
##      Chisq Df Pr(>Chisq)
## (Intercept) 28.543  1  9.163e-08 ***
## SI          82.959  1  < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
r.squaredGLMM(sUASndvi_yield_mixlm)

##      R2m      R2c
## [1,] 0.5352913 0.8488972

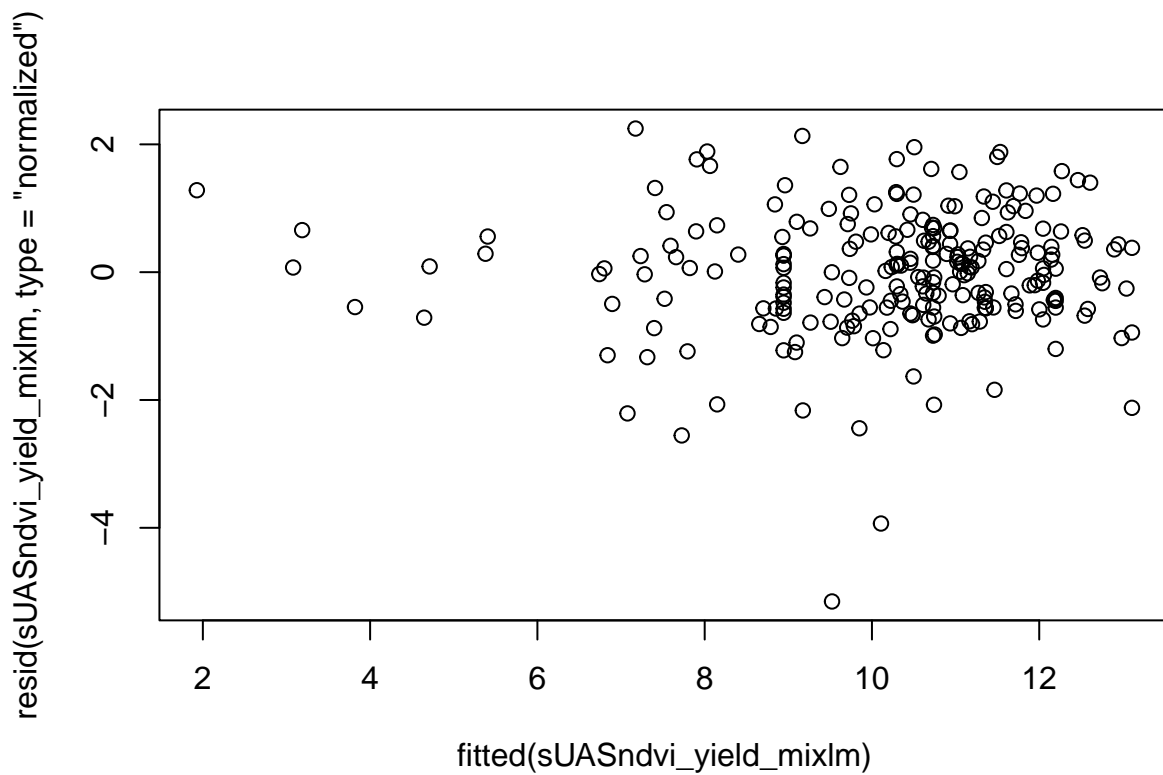
plot (sUASndvi_yield_mixlm)

```



diagnostics

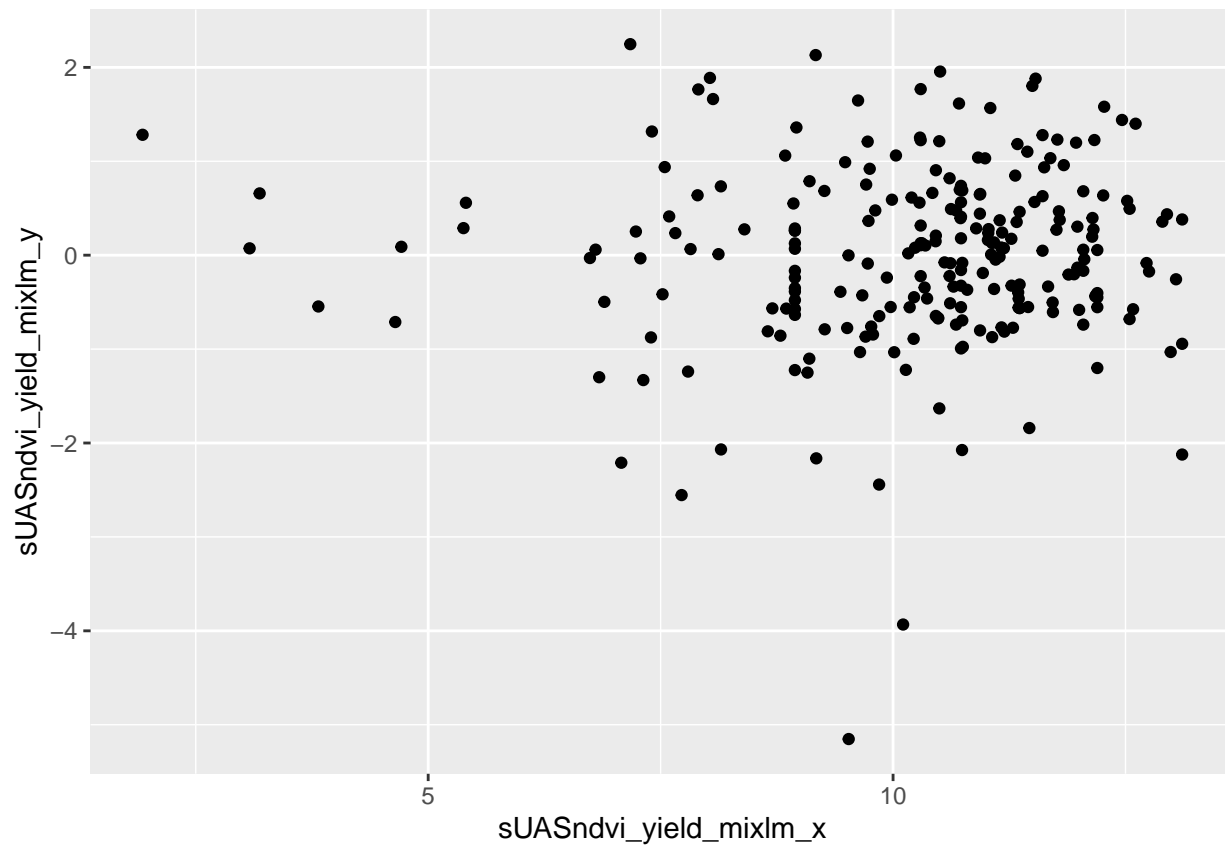
```
plot(resid(sUASndvi_yield_mixlm, type = "normalized") ~fitted(sUASndvi_yield_mixlm))
```



```
sUASndvi_yield_mixlm_y <- resid(sUASndvi_yield_mixlm, type = "normalized")
sUASndvi_yield_mixlm_x <- fitted(sUASndvi_yield_mixlm)
```

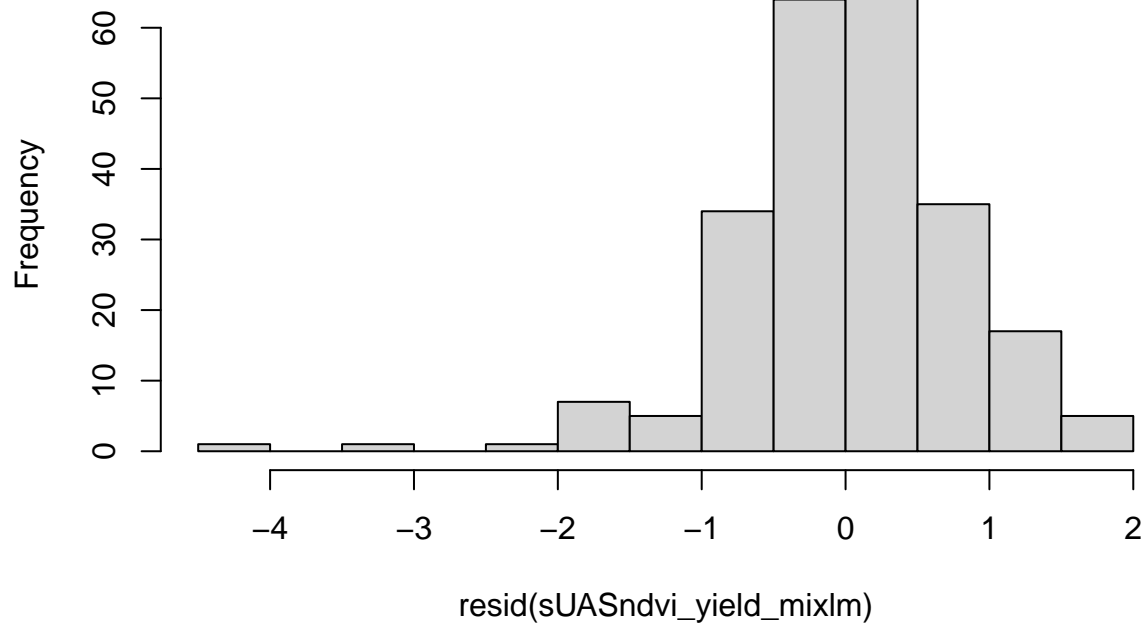
```
sUASndvi_yield_mixlmresid_data <- data.frame(sUASndvi_yield_mixlm_x , sUASndvi_yield_mixlm_y)
```

```
ggplot( data = sUASndvi_yield_mixlmresid_data , aes( x = sUASndvi_yield_mixlm_x , y = sUASndvi_yield_mi  
geom_point(mapping = aes(sUASndvi_yield_mixlm_x , sUASndvi_yield_mixlm_y) , data = sUASndvi_yield_mix
```

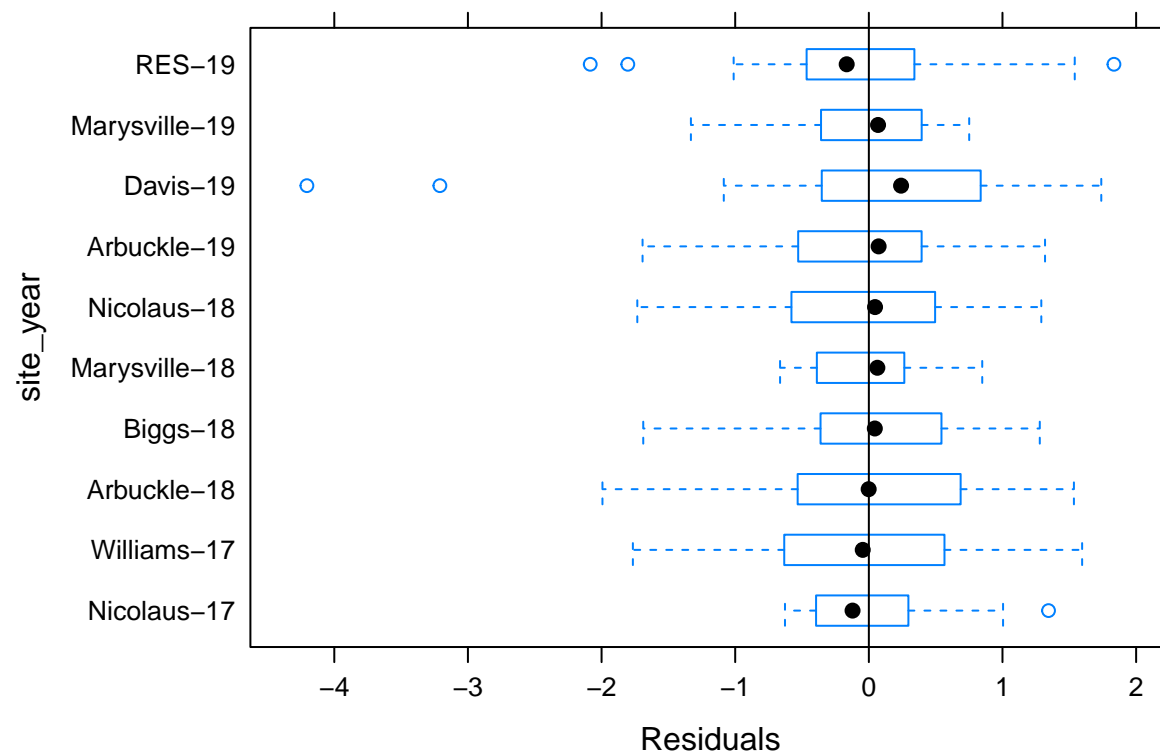


```
hist(resid(sUASndvi_yield_mixlm))
```

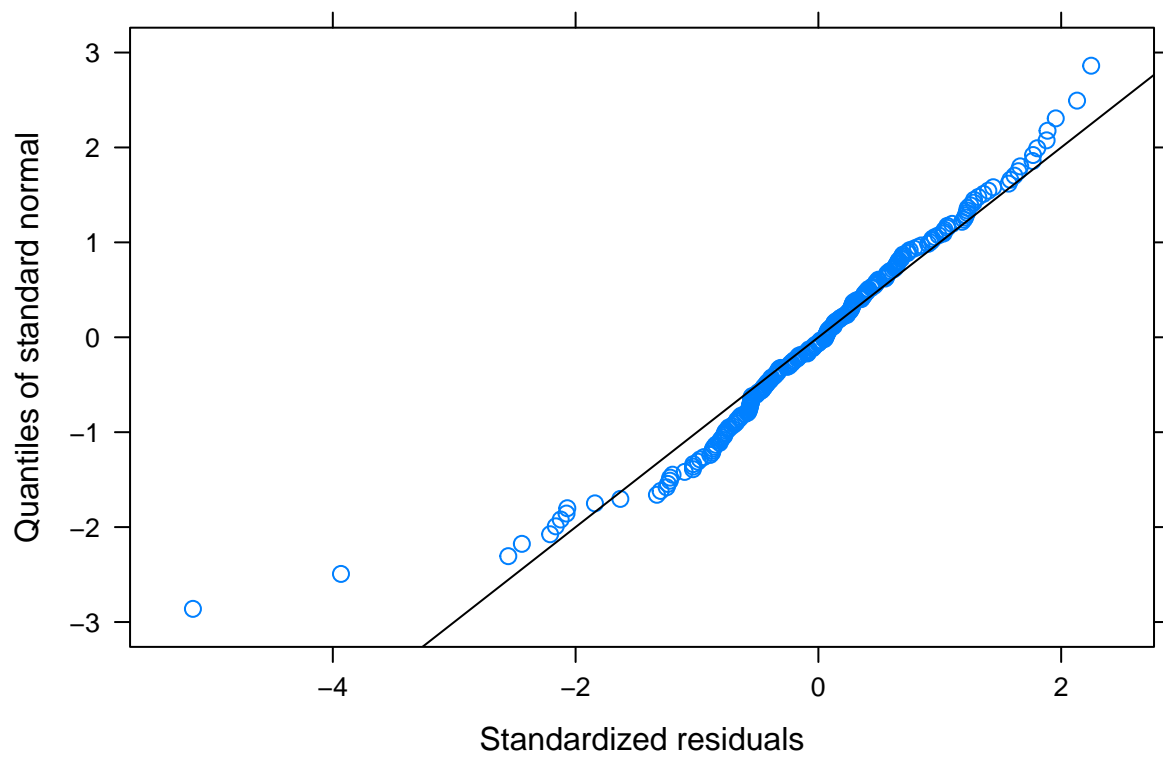

Histogram of resid(sUASndvi_yield_mixlm)



```
plot(sUASndvi_yield_mixlm, site_year ~ resid(.), abline = 0)
```

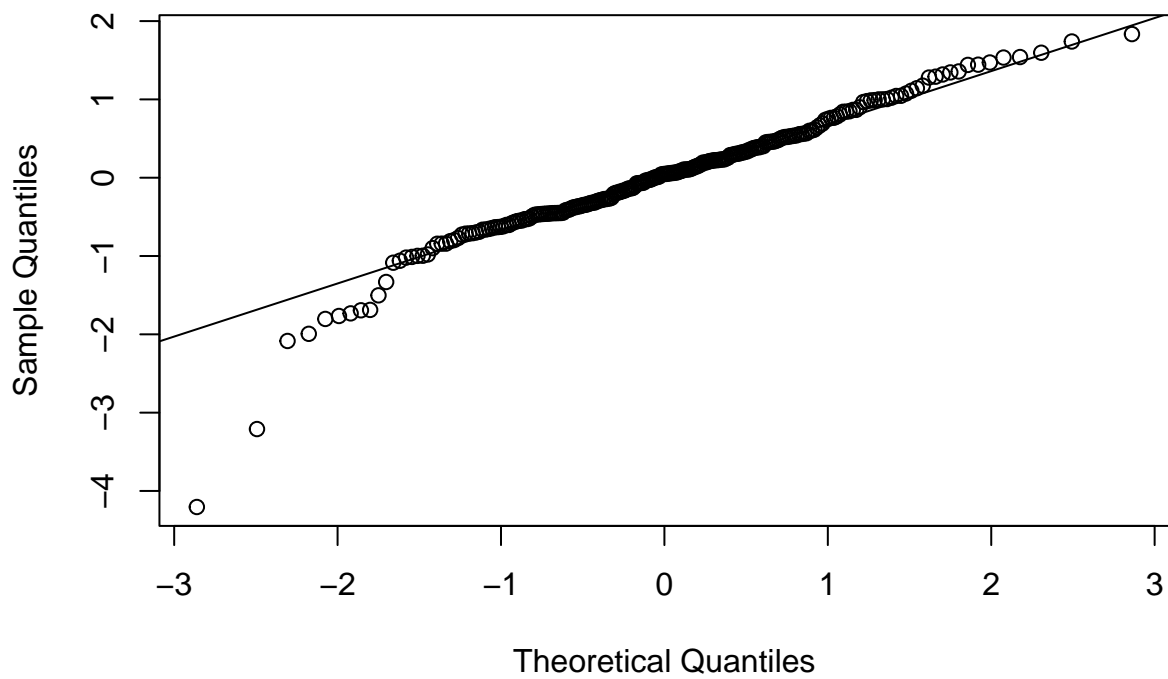


```
qqnorm(sUASndvi_yield_mixlm, abline = c(0,1) )
```

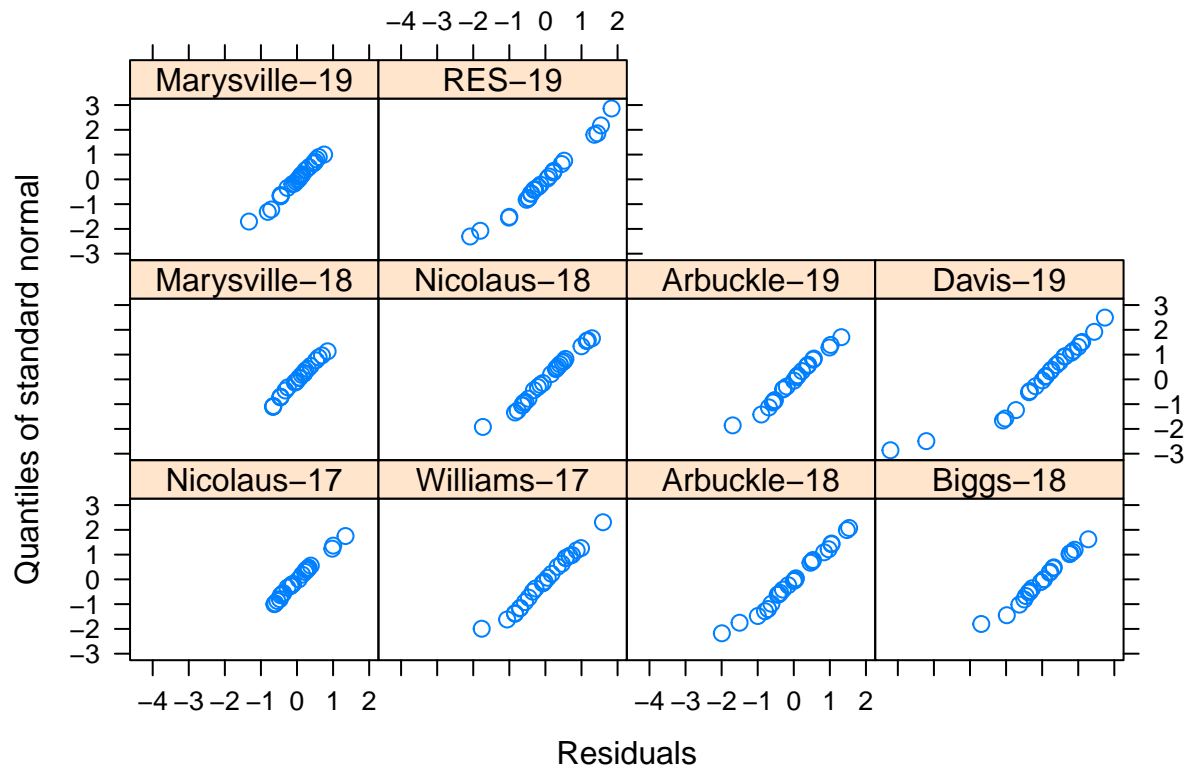


```
qqnorm(resid(sUASndvi_yield_mixlm))
qqline(resid(sUASndvi_yield_mixlm))
```

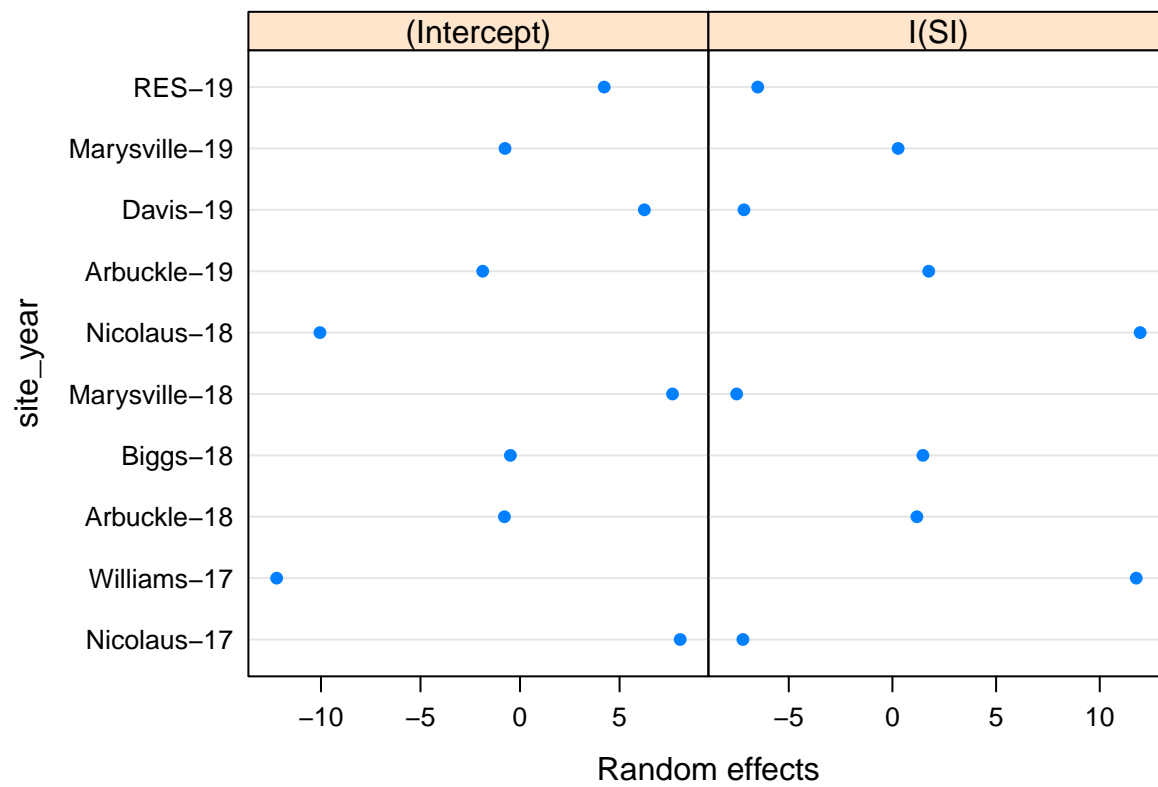
Normal Q–Q Plot



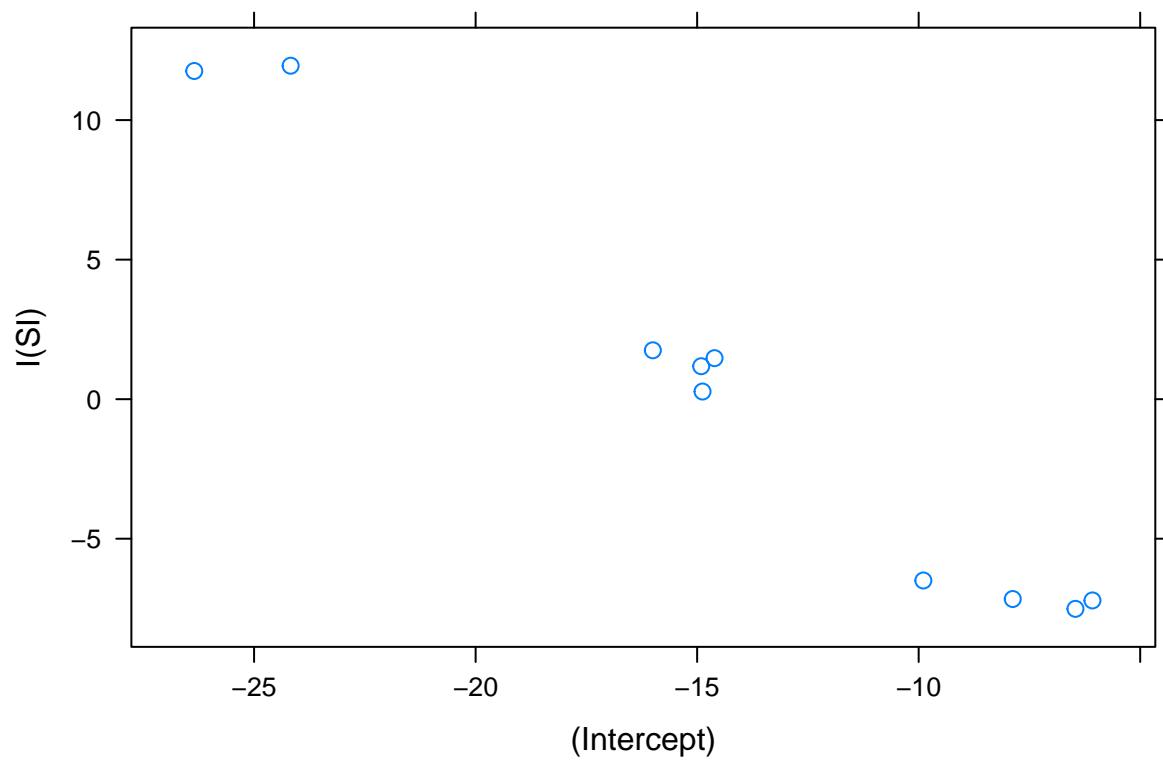
```
qqnorm(sUASndvi_yield_mixlm , ~resid(.) | site_year)
```



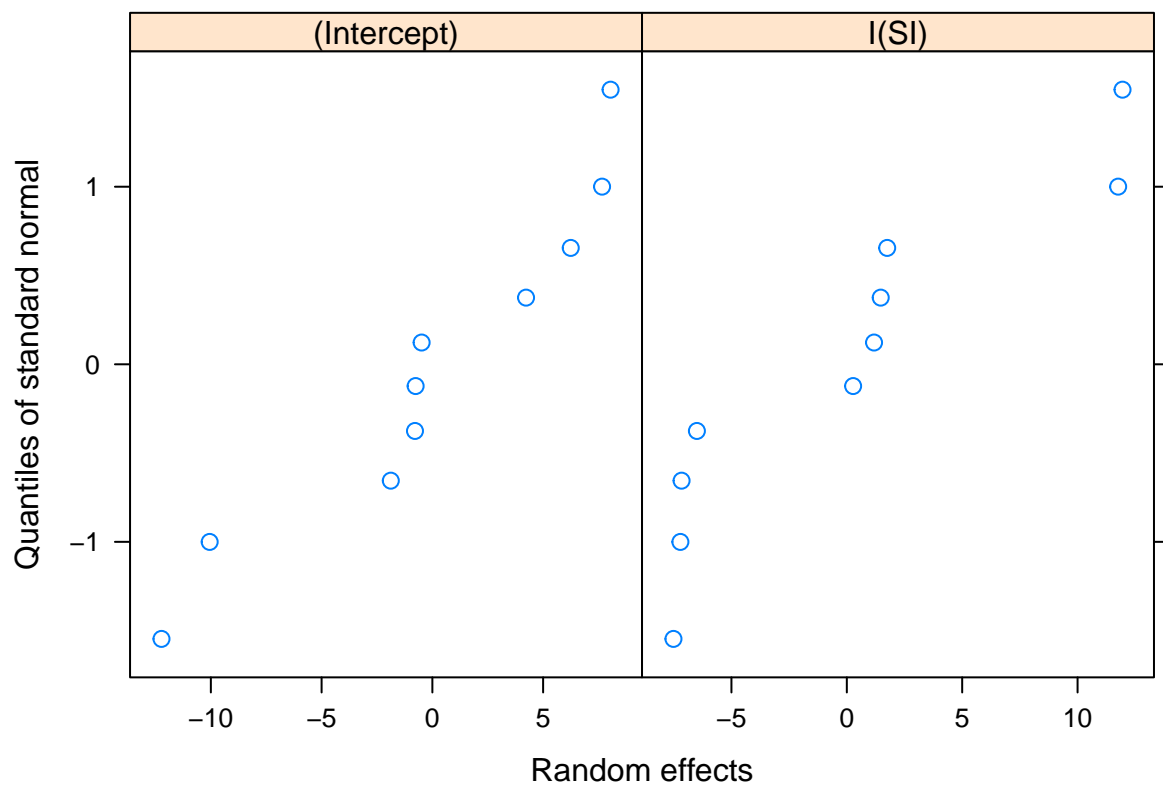
```
plot(ranef(sUASndvi_yield_mixlm))
```



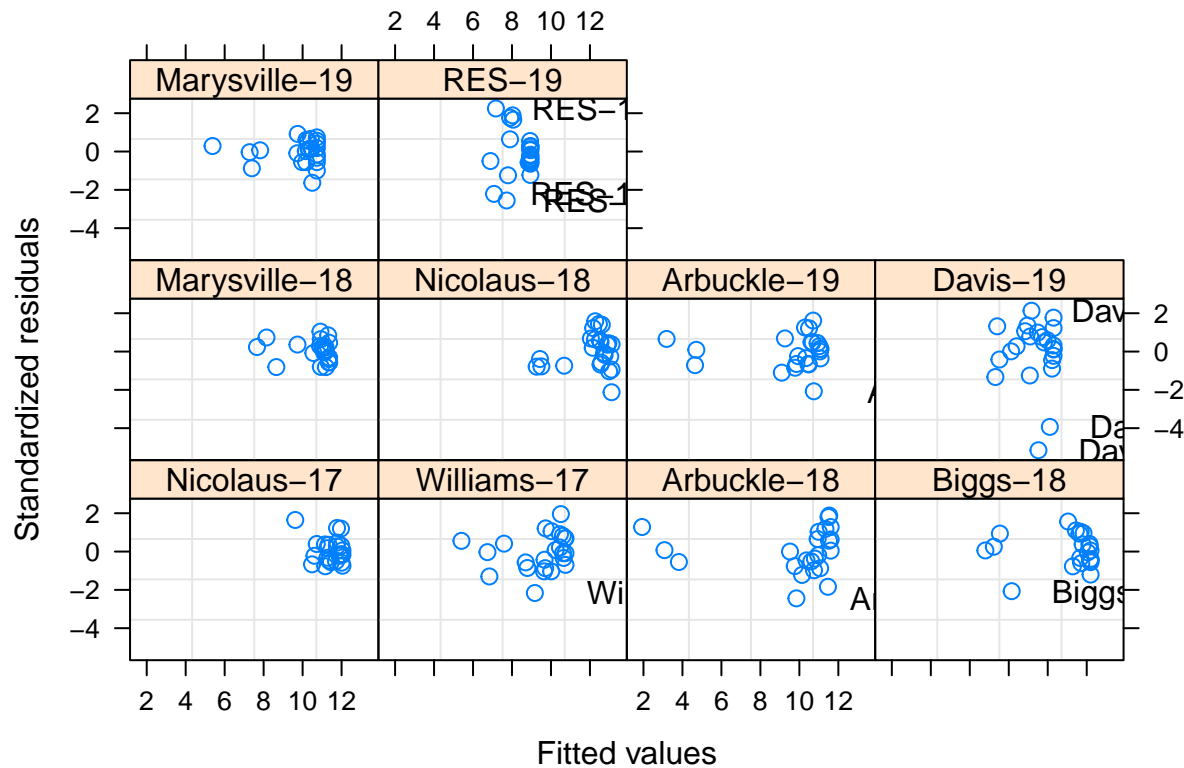
```
pairs(sUASndvi_yield_mixlm , id = 0.1)
```



```
qqnorm(sUASndvi_yield_mixlm , ~ranef(.))
```



```
plot( sUASndvi_yield_mixlm, resid(., type = "p") ~ fitted(.) | site_year,
      id = 0.05, adj = -0.3 )
```



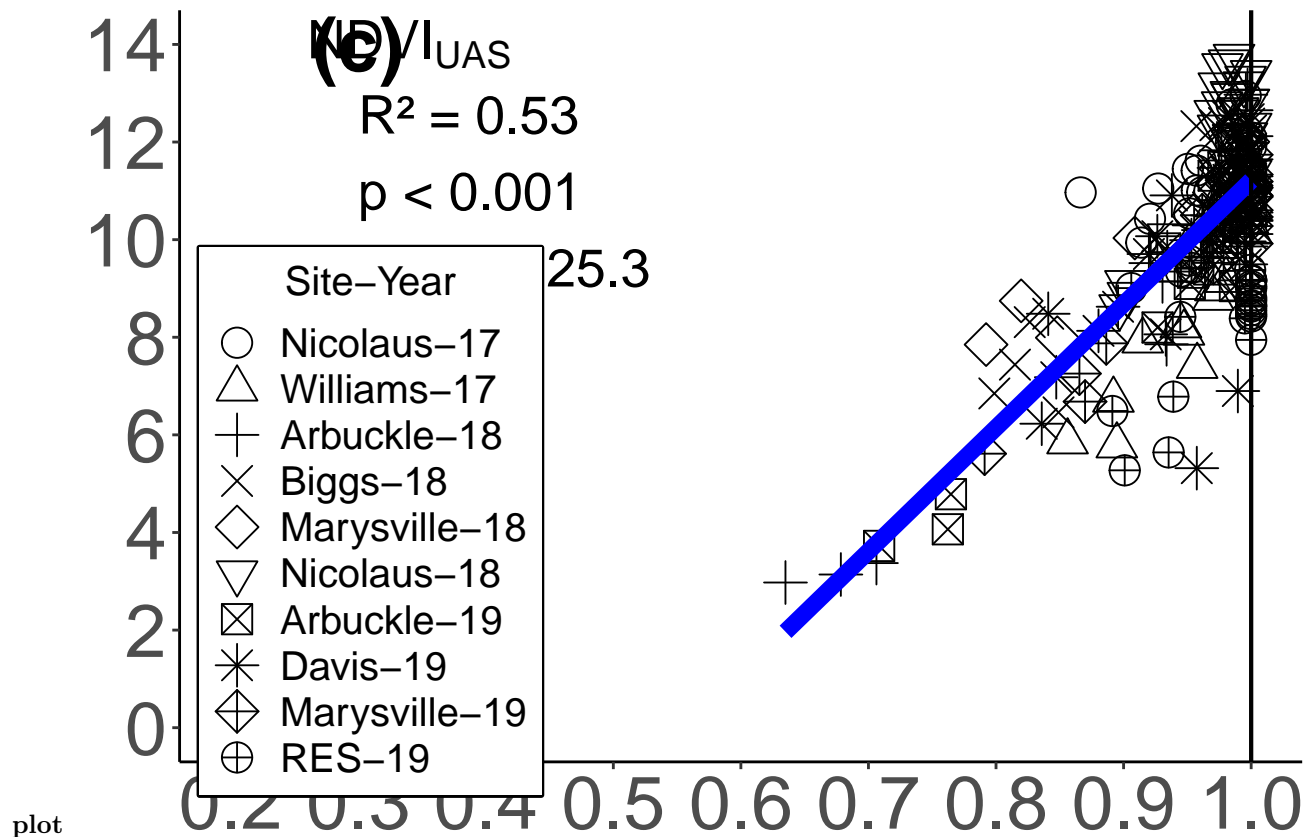
```
mylist <- list(SI=seq(round(min(paper3_uas_ndvi_data$SI), digits = 3) , 1 , by = .001))

sUASndvi_emmeans <- as.data.frame(summary(emmeans(sUASndvi_yield_mixlm , ~ SI , at = mylist )))
```

emmeans

```
plot_si2 <- ggplot( data = paper3_uas_ndvi_data , aes( x = SI , y = GrainYield_Mgha)) +
  geom_point( data = paper3_uas_ndvi_data , aes( x = SI , y = GrainYield_Mgha , shape = site_year) , size = 3) +
  geom_line( data = sUASndvi_emmeans , aes(x = SI , y = emmean) , color = "blue" , size = 3) +
  theme_classic() +
  labs( x = NULL , y = NULL , shape = "Site-Year") +
  scale_x_continuous(breaks = seq(0.2 , 1 , by = .1)) +
  scale_y_continuous(breaks = seq(0 , 14 , by = 2)) +
  coord_cartesian(ylim = c(0,14) , xlim = c(0.2 , 1)) +
  geom_vline(xintercept = 1 , size = 0.75) +
  theme(axis.text = element_text(size = 28),
        axis.title = element_text(size = 28),
        legend.text = element_text(size = 15),
        legend.title = element_text(size = 15 , hjust = 0.5),
        legend.position = c(.17 , .32),
        legend.box.background = element_rect(size = 1)) +
  annotate("text", x=0.3, y=11, label="R² = 0.53\np < 0.001\nslope = 25.3", size=7 , color="black" , hjust = 0.5 , fontface = 2) +
  annotate("text", x=0.3, y=14, label="(c)", size=10 , color = "black" , hjust = 0.5 , fontface = 2) +
  annotate("text", x=0.34, y=14, label="NDVI[UAS]", size=7 , color = "black" , hjust = 0.5 , parse = TRUE) +
  scale_shape_manual(values = c(1:20))
```

plot_si2



sUAS ndre RI

```

sUASndre_yield_mixlm <- lme(fixed = GrainYield_Mgha ~ SI ,
                             random = ~I(SI) | site_year,
                             data = paper3_uas_ndre_data)

```

```
summary(sUASndre_yield_mixlm)
```

model

```

## Linear mixed-effects model fit by REML
##   Data: paper3_uas_ndre_data
##       AIC      BIC    logLik
##   603.415 624.1725 -295.7075
##
## Random effects:
##   Formula: ~I(SI) | site_year
##   Structure: General positive-definite, Log-Cholesky parametrization
##              StdDev    Corr
## (Intercept) 3.5144210 (Intr)
## I(SI)        3.6847042 -0.946
## Residual    0.7451118
##
## Fixed effects: GrainYield_Mgha ~ SI
##              Value Std.Error   DF   t-value p-value
## (Intercept) -0.21627  1.176285  226  -0.183859  0.8543
## SI           11.74458  1.240542  226   9.467298  0.0000

```

```
## Correlation:
## (Intr)
## SI -0.951
##
## Standardized Within-Group Residuals:
##      Min      Q1      Med      Q3      Max
## -5.096822800 -0.549578194  0.001105708  0.517734896  2.092222985
##
## Number of Observations: 237
## Number of Groups: 10
```

```
summary(sUASndre_yield_mixlm)$tTable
```

```
##              Value Std.Error  DF    t-value      p-value
## (Intercept) -0.2162704  1.176285 226 -0.1838588 8.542891e-01
## SI          11.7445785  1.240542 226  9.4672982 3.983273e-18
```

```
Anova(sUASndre_yield_mixlm , type = 2)
```

```
## Analysis of Deviance Table (Type II tests)
##
## Response: GrainYield_Mgha
##      Chisq Df Pr(>Chisq)
## SI 89.63  1  < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

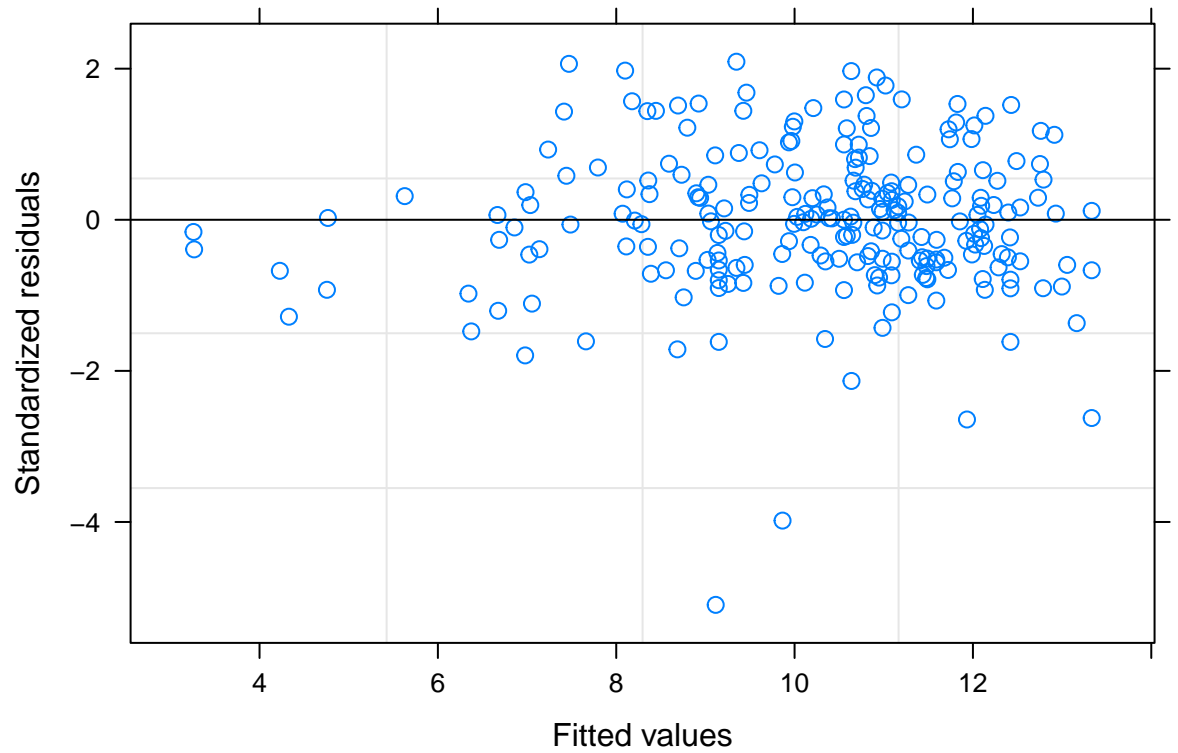
```
Anova(sUASndre_yield_mixlm , type = 3)
```

```
## Analysis of Deviance Table (Type III tests)
##
## Response: GrainYield_Mgha
##              Chisq Df Pr(>Chisq)
## (Intercept)  0.0338  1    0.8541
## SI          89.6297  1    <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
r.squaredGLMM(sUASndre_yield_mixlm)
```

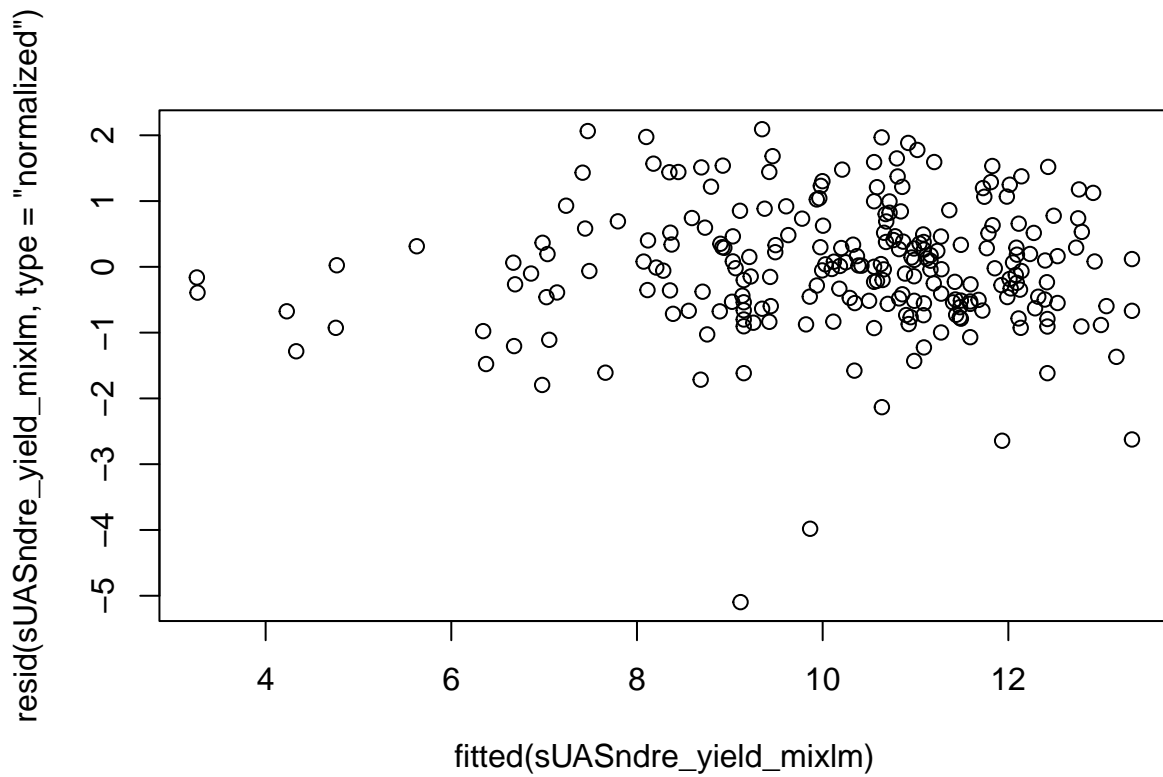
```
##              R2m      R2c
## [1,] 0.5075114 0.8677391
```

```
plot (sUASndre_yield_mixlm)
```



diagnostics

```
plot(resid(sUASndre_yield_mixlm, type = "normalized") ~fitted(sUASndre_yield_mixlm))
```

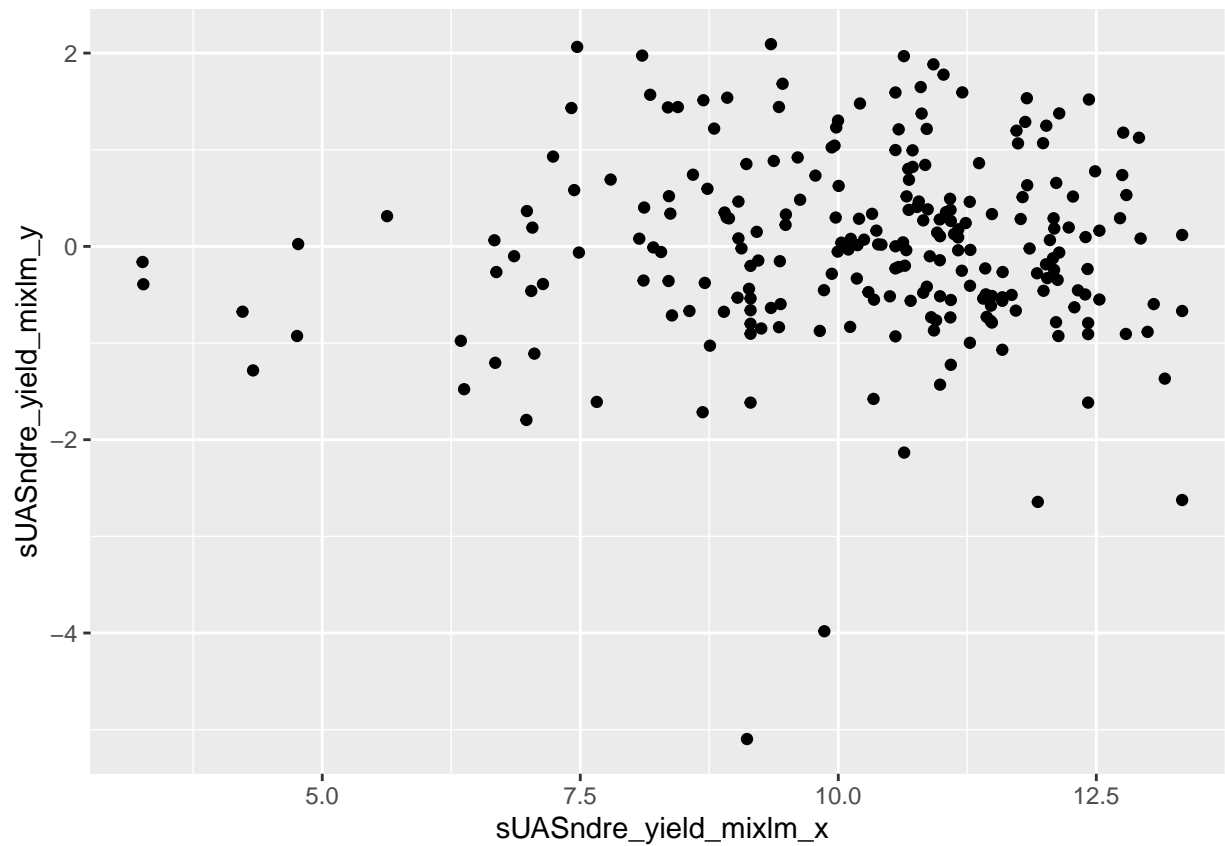


```
sUASndre_yield_mixlm_y <- resid(sUASndre_yield_mixlm, type = "normalized")
sUASndre_yield_mixlm_x <- fitted(sUASndre_yield_mixlm)
```



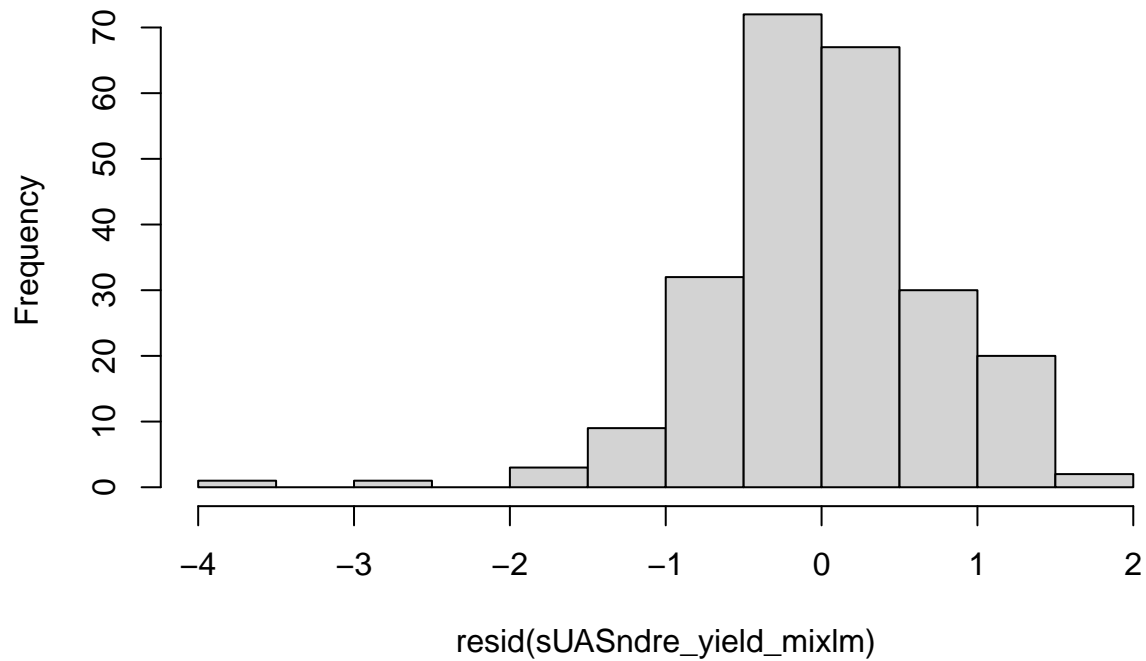
```
sUASndre_yield_mixlmresid_data <- data.frame(sUASndre_yield_mixlm_x , sUASndre_yield_mixlm_y)
```

```
ggplot( data = sUASndre_yield_mixlmresid_data , aes( x = sUASndre_yield_mixlm_x , y = sUASndre_yield_mi  
geom_point(mapping = aes(sUASndre_yield_mixlm_x , sUASndre_yield_mixlm_y) , data = sUASndre_yield_mix
```

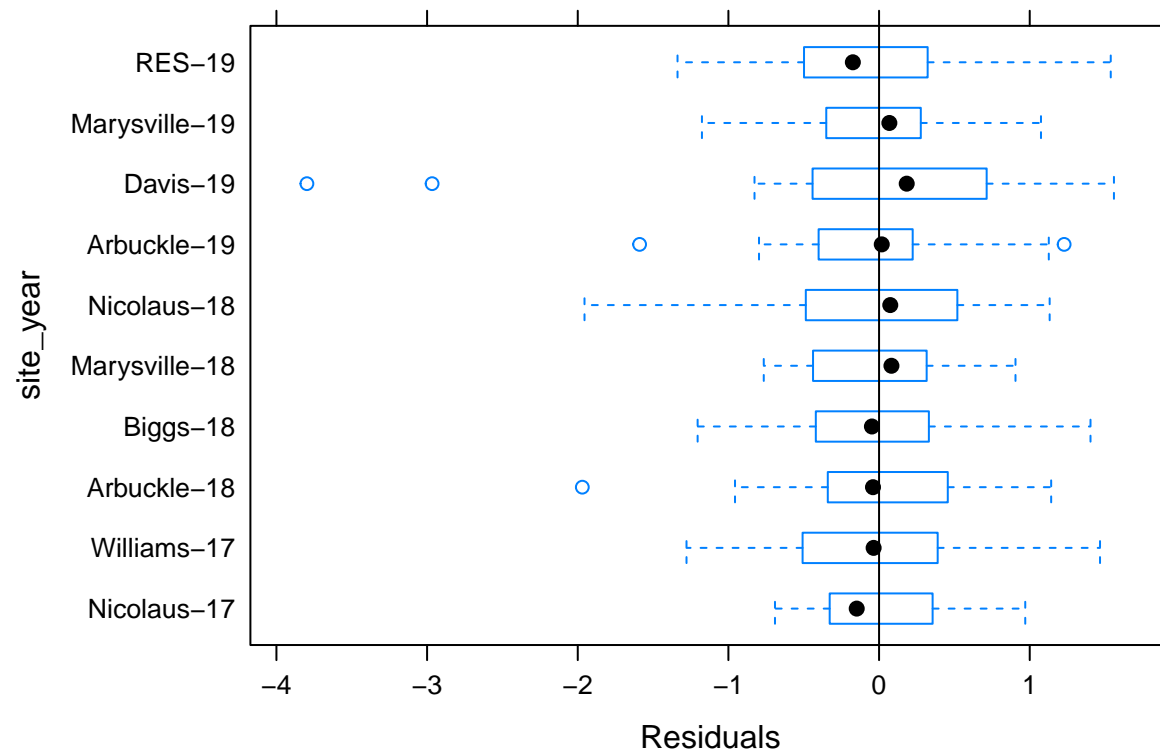


```
hist(resid(sUASndre_yield_mixlm))
```

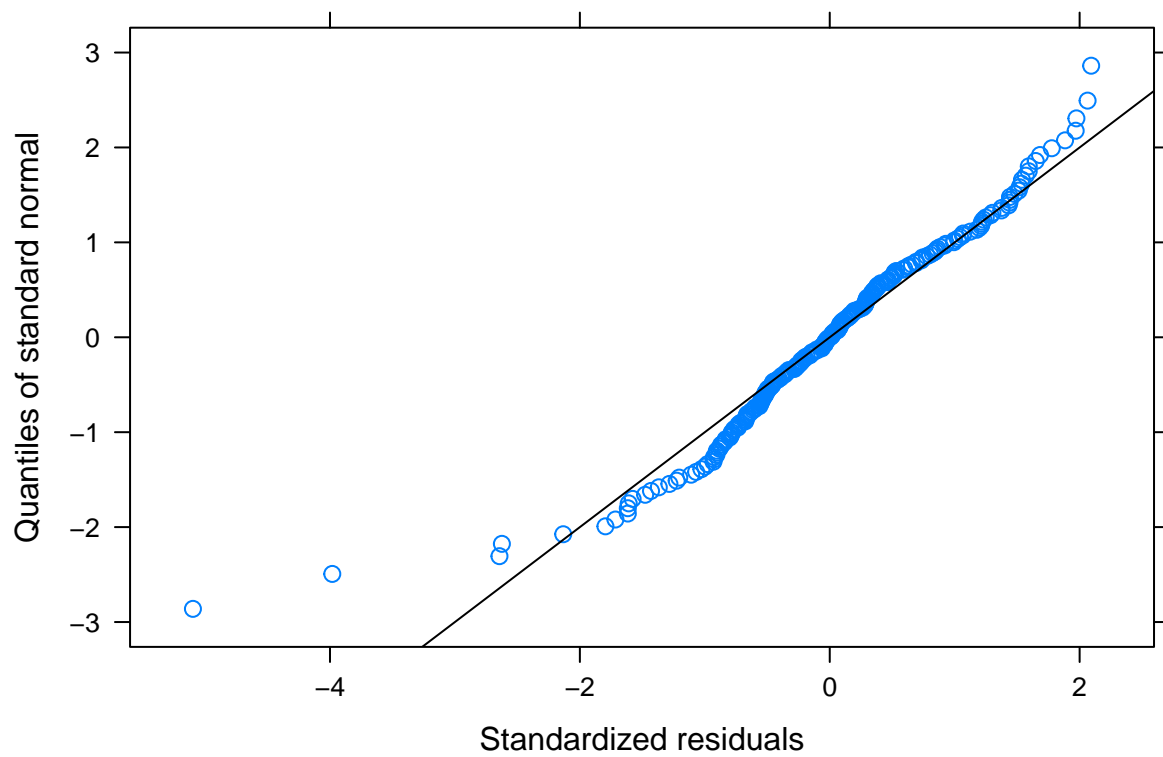
Histogram of resid(sUASndre_yield_mixlm)



```
plot(sUASndre_yield_mixlm, site_year ~ resid(.), abline = 0)
```

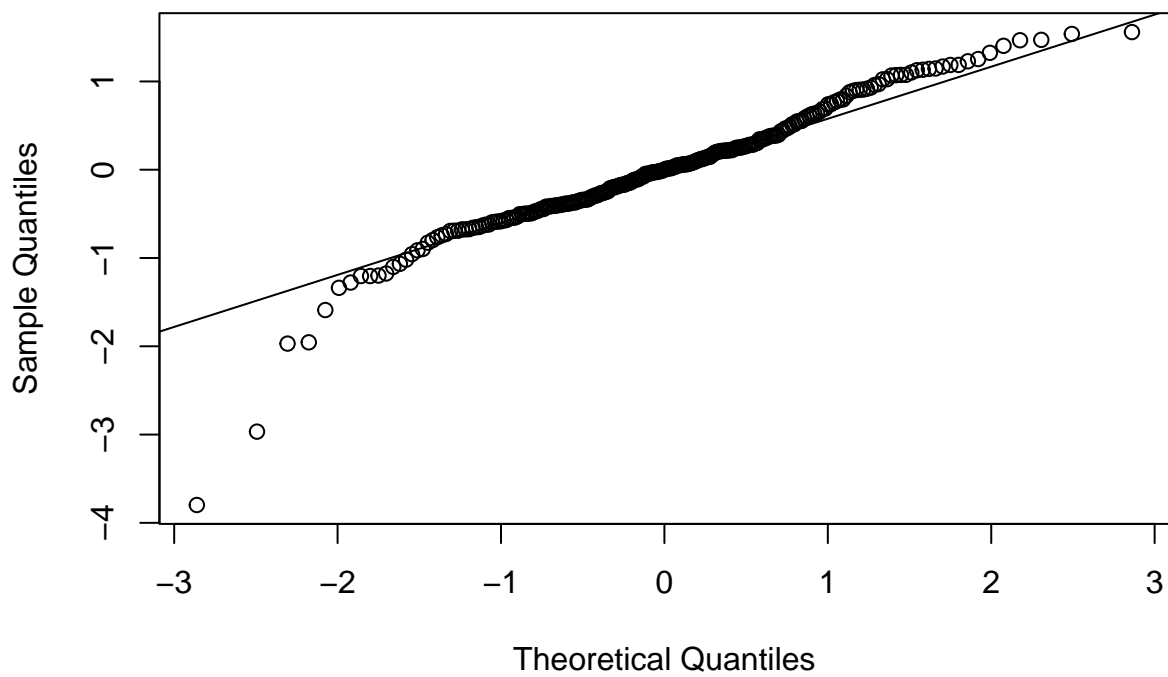


```
qqnorm(sUASndre_yield_mixlm, abline = c(0,1) )
```

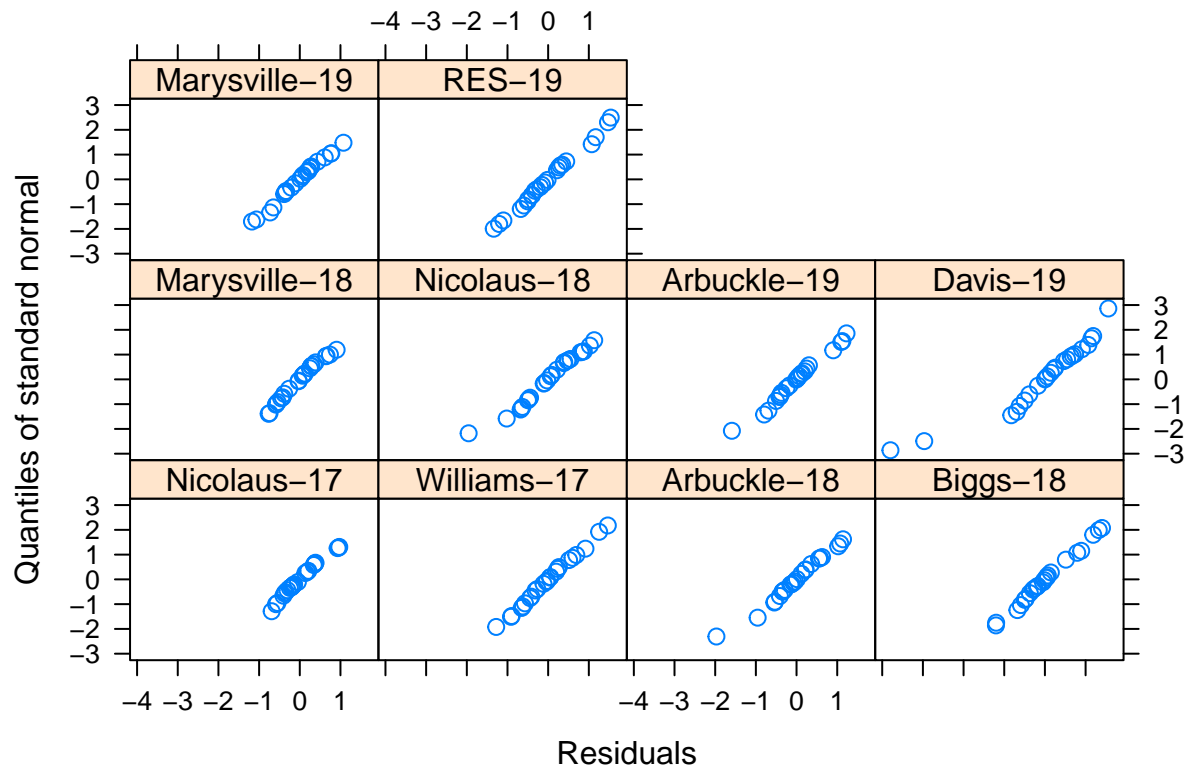


```
qqnorm(resid(sUASndre_yield_mixlm))
qqline(resid(sUASndre_yield_mixlm))
```

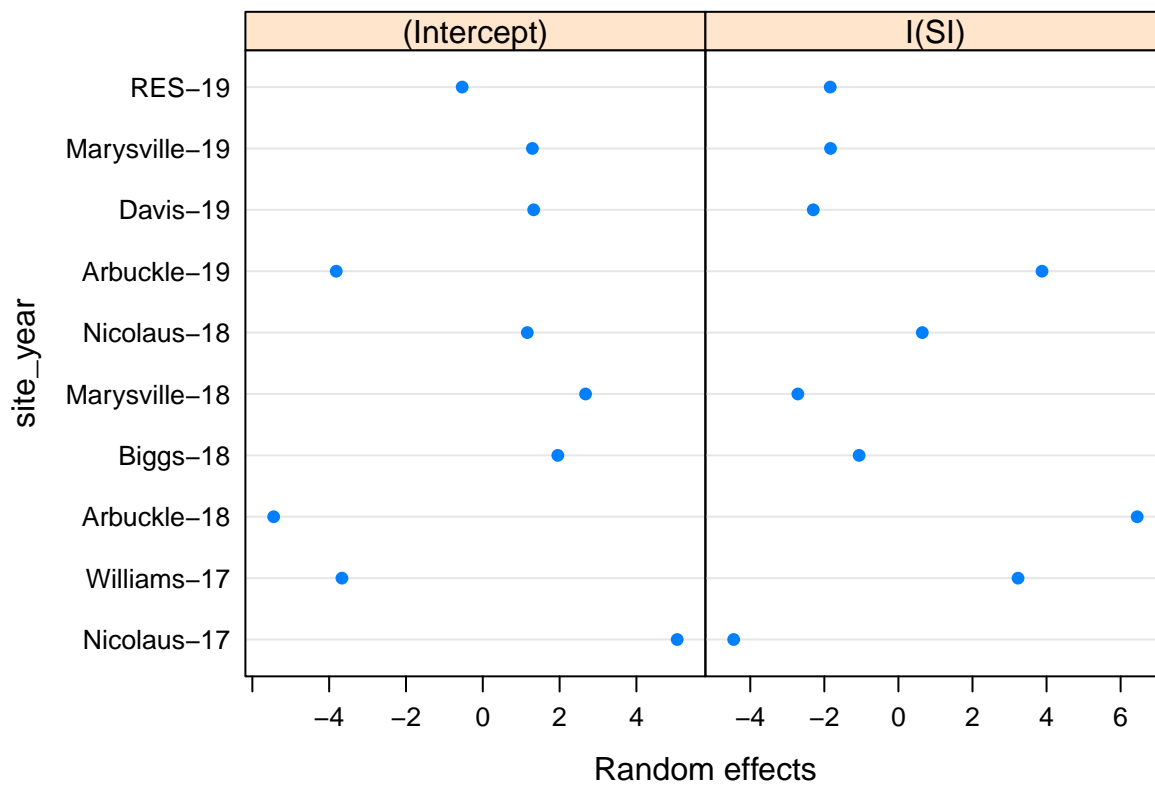
Normal Q–Q Plot



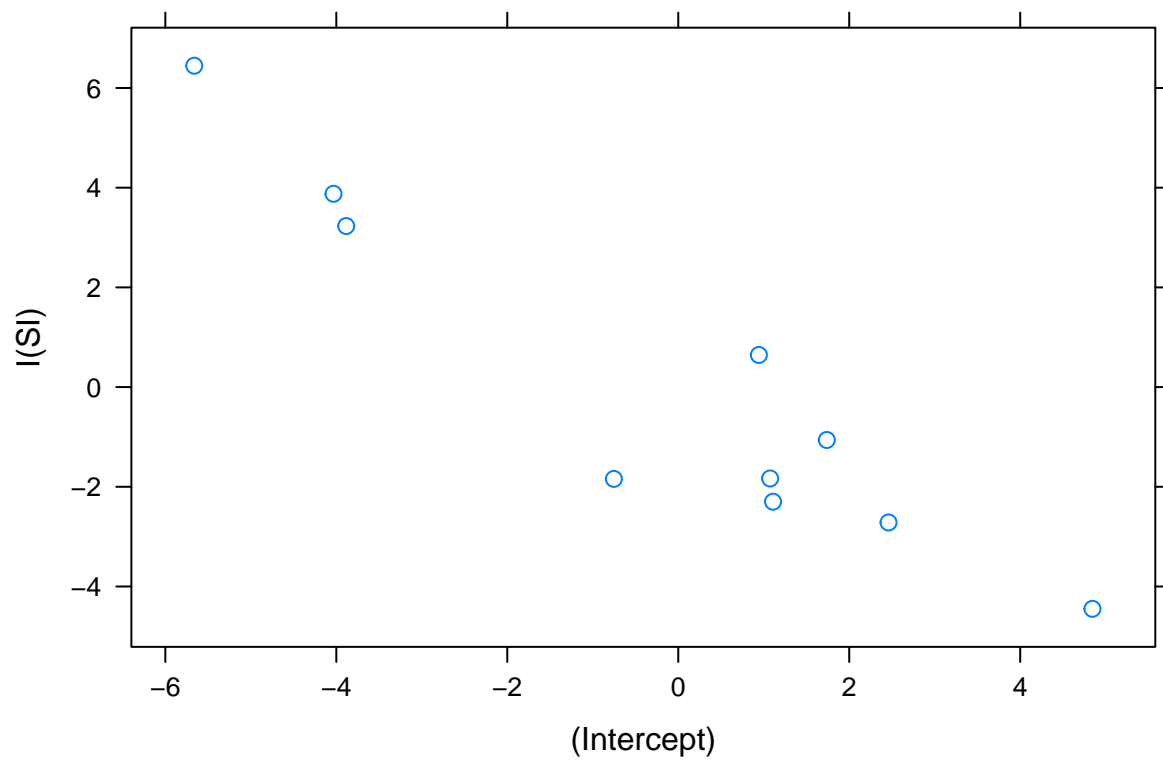
```
qqnorm(sUASndre_yield_mixlm , ~resid(.) | site_year)
```



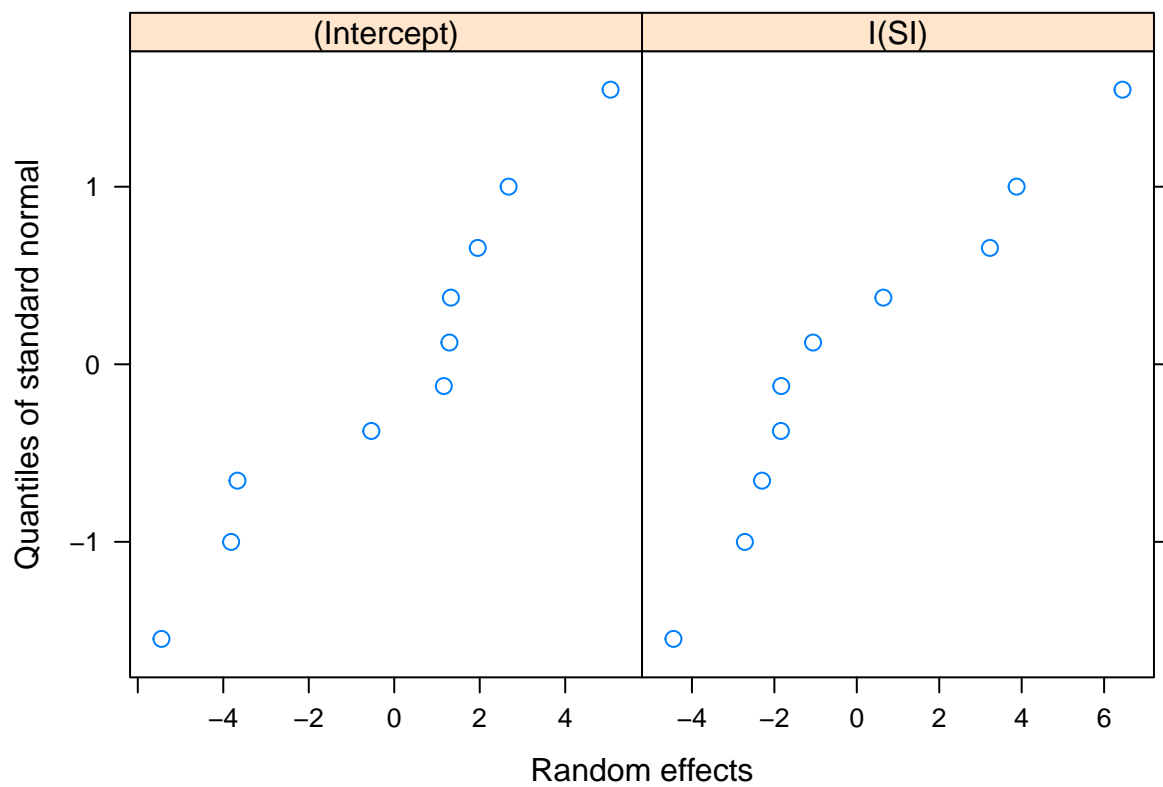
```
plot(ranef(sUASndre_yield_mixlm))
```



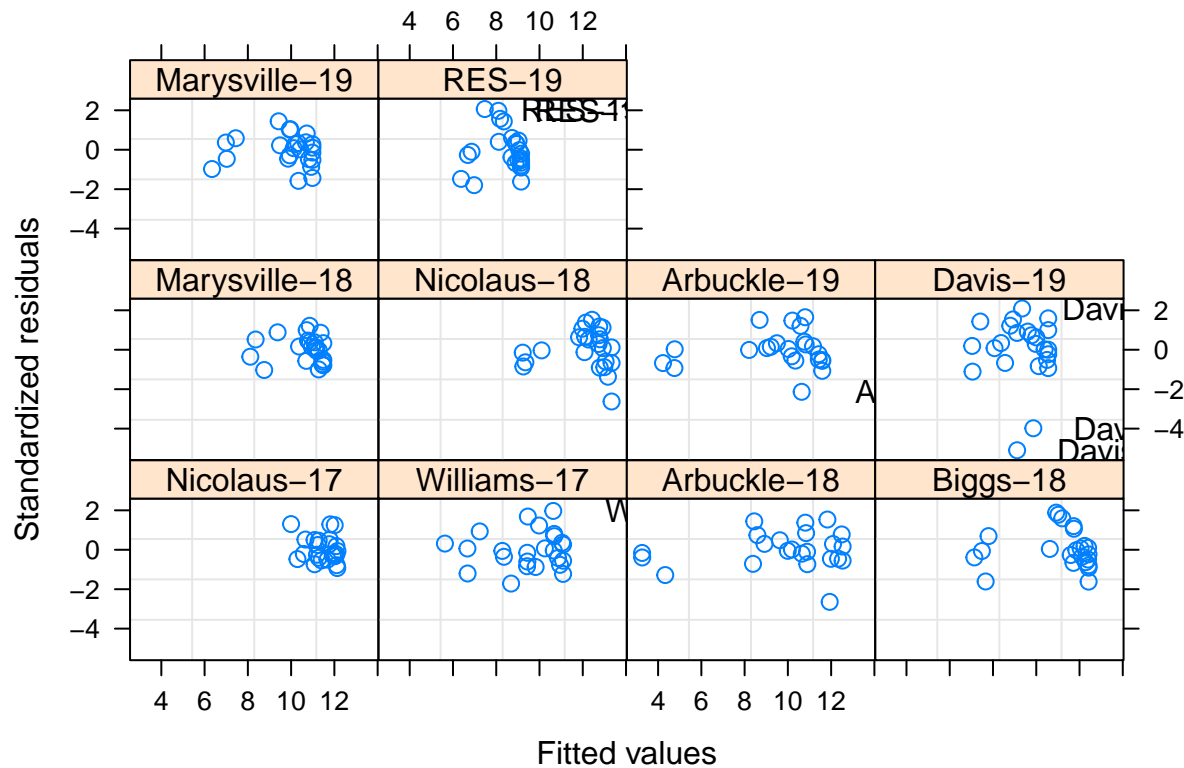
```
pairs(sUASndre_yield_mixlm , id = 0.1)
```



```
qqnorm(sUASndre_yield_mixlm , ~ranef(.))
```



```
plot( sUASndre_yield_mixlm, resid(., type = "p") ~ fitted(.) | site_year,
      id = 0.05, adj = -0.3 )
```



```
mylist <- list(SI=seq(round(min(paper3_uas_ndre_data$SI), digits = 3) , 1 , by = .001))

sUASndre_emmeans <- as.data.frame(summary(emmeans(sUASndre_yield_mixlm , ~ SI , at = mylist )))
```

emmeans

```
plot_si3 <- ggplot( data = paper3_uas_ndre_data , aes( x = SI , y = GrainYield_Mgha)) +
  geom_point( data = paper3_uas_ndre_data , aes( x = SI , y = GrainYield_Mgha , shape = site_year) , size = 3) +
  geom_line( data = sUASndre_emmeans , aes(x = SI , y = emmean) , color = "blue" , size = 3) +
  theme_classic() +
  labs( x = NULL , y = NULL , shape = "Site-Year") +
  scale_x_continuous(breaks = seq(0.2 , 1 , by = .1)) +
  scale_y_continuous(breaks = seq(0 , 14 , by = 2)) +
  coord_cartesian(ylim = c(0,14) , xlim = c(0.2 , 1)) +
  geom_vline(xintercept = 1 , size = 0.5) +
  theme(axis.text = element_text(size = 28),
        axis.title = element_text(size = 28),
        legend.text = element_text(size = 20),
        legend.title = element_text(size = 20),
        legend.position = "none") +
  annotate("text", x=0.3, y=11, label="R² = 0.51\np < 0.001\nslope = 11.7", size=7 , color = "black" , fontface = "italic") +
  annotate("text", x=0.3, y=14, label="(b)", size=10 , color = "black" , hjust = 0.5 , fontface = 2) +
  annotate("text", x=0.34, y=14, label="NDRE[UAS]", size=7 , color = "black" , hjust = 0 , parse = T) +
  scale_shape_manual(values = c(1:20))

plot_si3
```

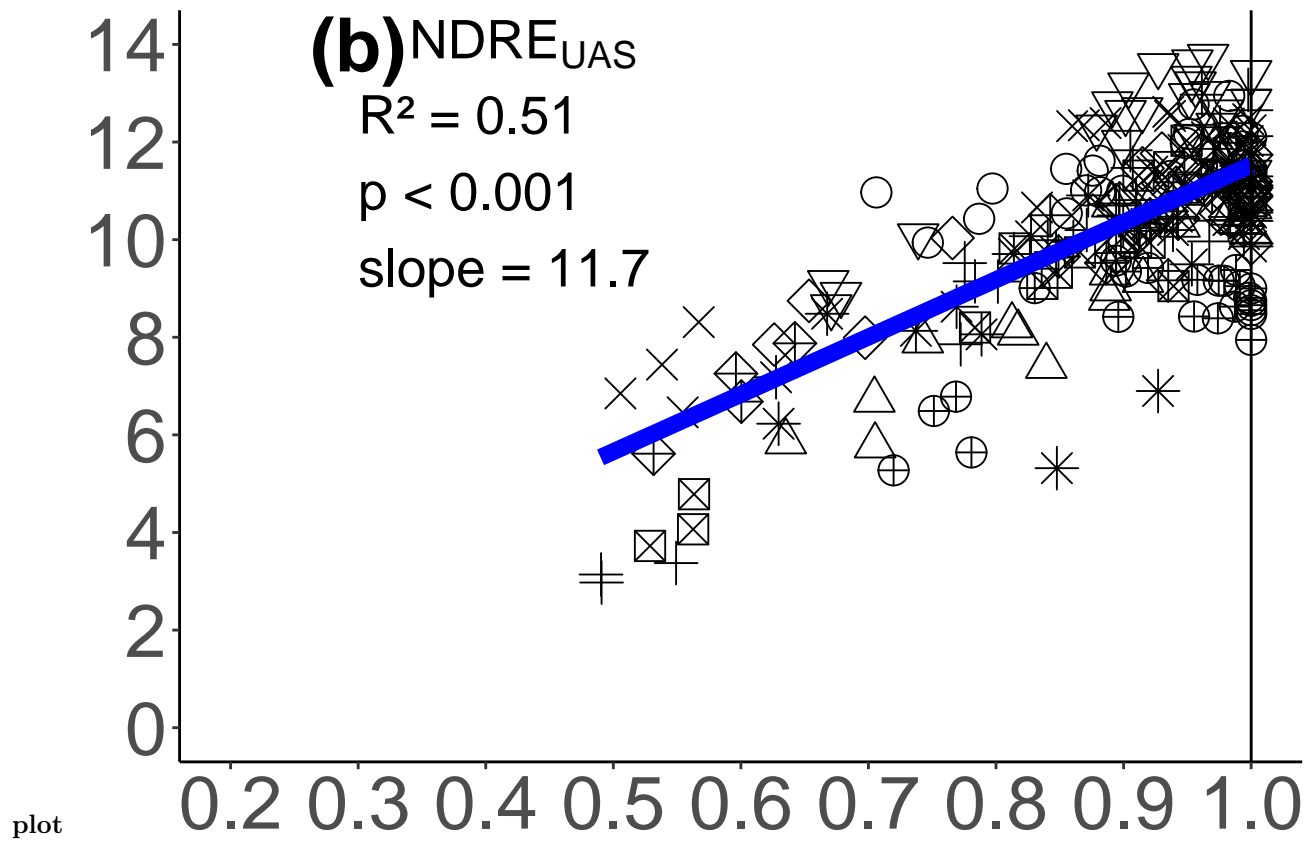
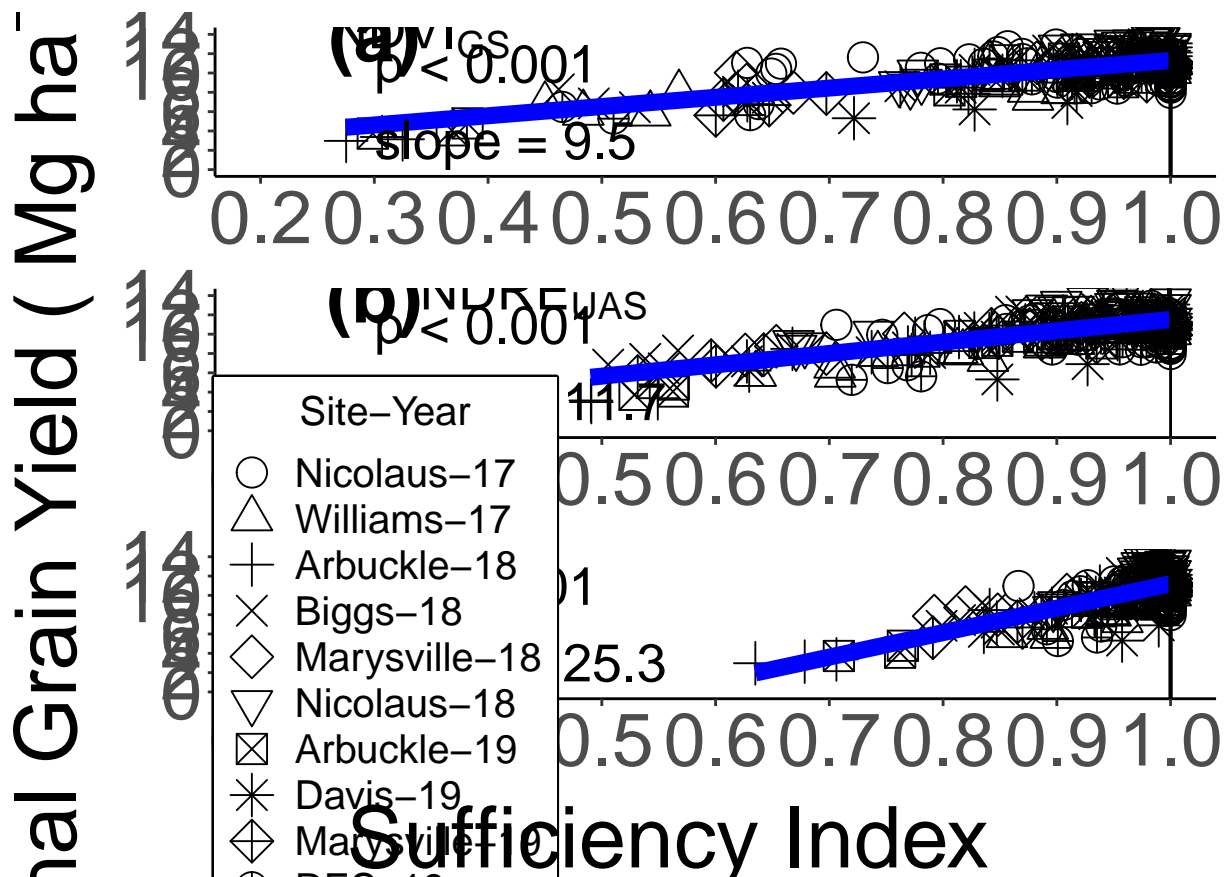


FIGURE 5

```
Fig5 <- grid.arrange(arrangeGrob(plot_si1,
                                  plot_si3,
                                  plot_si2,
                                  nrow = 3,
                                  ncol = 1,
                                  left = textGrob("Final Grain Yield ( Mg ha-1~") , rot = 90 , gp = gpar(fontsize = 32)),
                                  bottom = textGrob("Sufficiency Index" , gp = gpar(fontsize = 32))))
```



ggsave("FIGURES/Fig5.tiff" , Fig5 , compression = "lzw" , width = 22 , height = 22, type = "cairo" , dp