

Untitled3

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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:

START

```
Sys.time()
```

```
## [1] "2022-02-21 16:02:29 PST"
```

```
library(tinytex)
library(ggplot2)
library(tidyverse)
library(cowplot)
library(Cairo)
library(modelr)
library(gridExtra)
library(mixtools)
library(nlme)
library(car)
library(emmeans)
library(MuMIn)
library(ggpmisc)
library(gridExtra)
library(gtable)
library(grid)
library(RColorBrewer)
library(segmented)
library(data.table)
library(scales)
```

```
#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[,17] [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 analysis and also the N rate tha
str(gs_ndvi_data , give.attr = FALSE)
```

```
## spec_tbl_df[,17] [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[,18] [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,

```

```

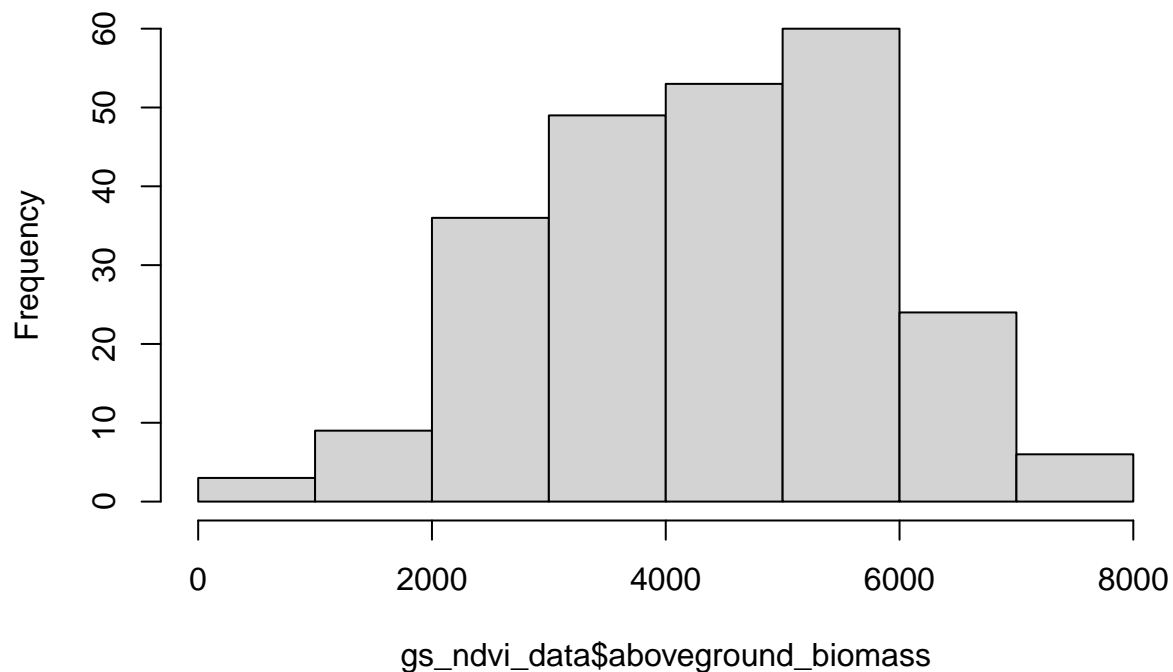
plot,
N_level_kgha,
aboveground_biomass,
n_content,
PI_N_Uptake,
NDVI) #selects the relevant columns

gs_ndvi_data$site_year <- factor(gs_ndvi_data$site_year , levels = c( "Nicolaus-17" , "Williams-17" , "
gs_ndvi_data$NDVI <- round(gs_ndvi_data$NDVI , digits = 2) #rounds to the precision of the instrument

hist(gs_ndvi_data$aboveground_biomass)

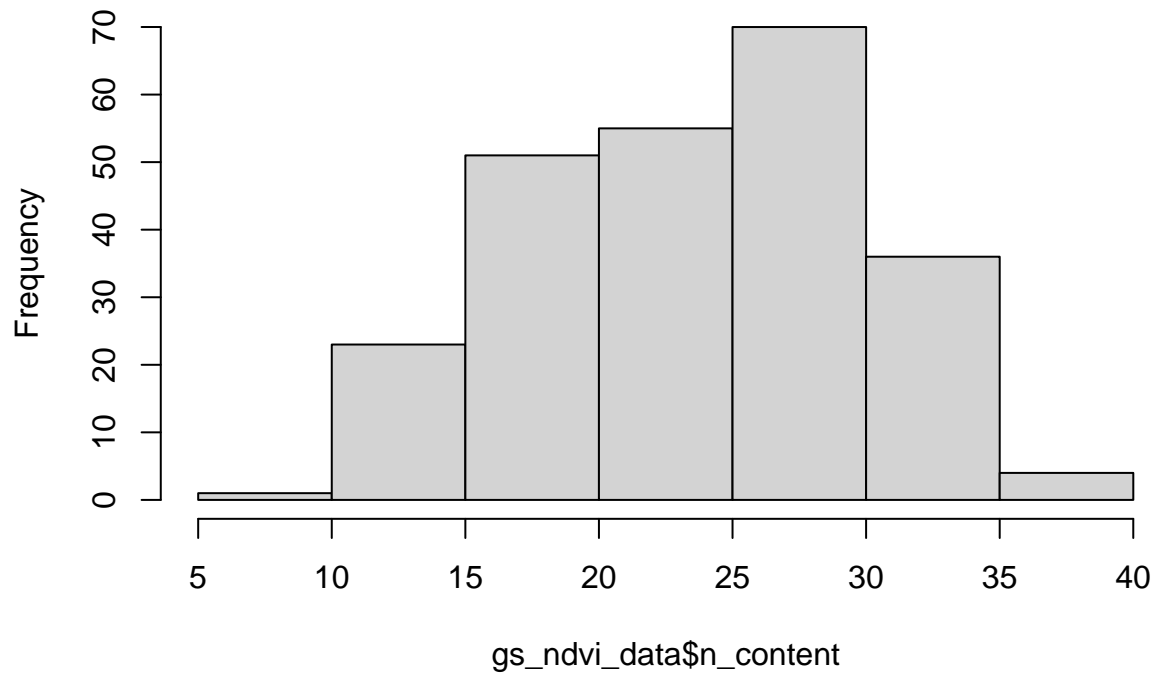
```

Histogram of gs_ndvi_data\$aboveground_biomass



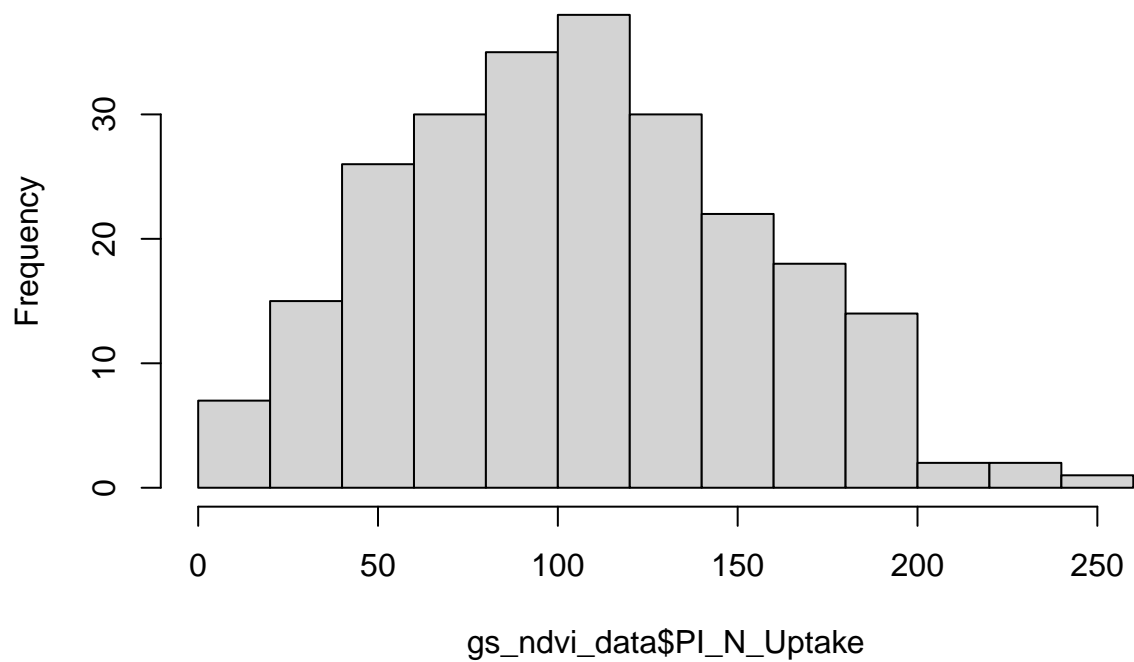
```
hist(gs_ndvi_data$n_content)
```

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[,34] [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"),
         N_level != "275") #removing the years and N rate to match with NDVI data
```

```

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)

```

```

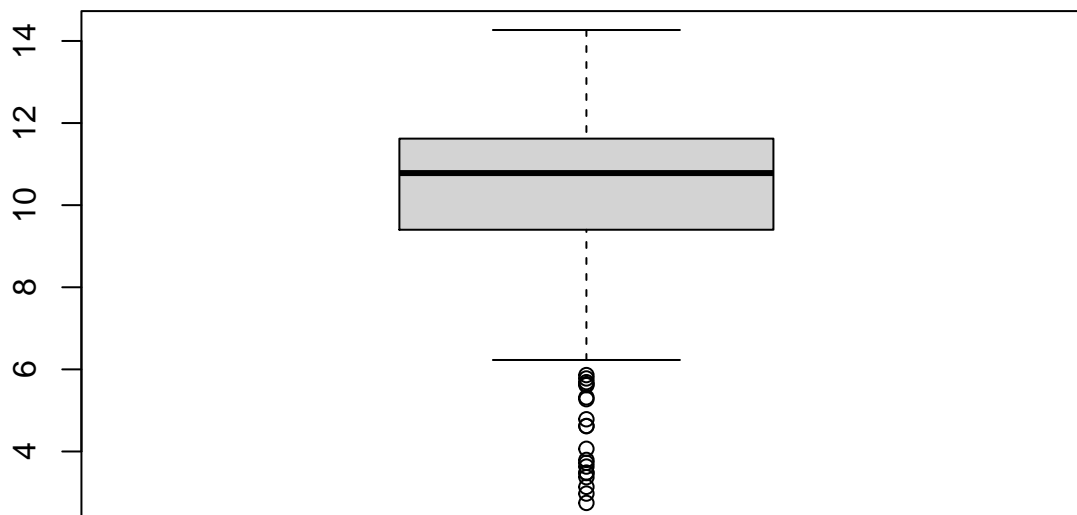
## spec_tbl_df[,38] [528 x 38] (S3: spec_tbl_df/tbl_df/tbl/data.frame)
## $ site_year      : Factor w/ 10 levels "Arbuckle-18",...: 7 7 7 7 7 7 7 7 7 7 ...
## $ year           : Factor w/ 3 levels "2017","2018",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ Block          : Factor w/ 4 levels "1","2","3","4": 1 1 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/ 4 levels "0","25","30",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ SeasonalNRate   : num [1:528] 225 0 125 175 45 75 75 125 0 175 ...
## $ N_level_kgha    : num [1:528] 225 0 125 175 45 75 75 125 0 175 ...
## $ TopDress_kgha   : num [1:528] 0 0 0 0 0 0 0 0 0 0 ...
## $ SeasonalNRate_kgha : num [1:528] 225 0 125 175 45 75 75 125 0 175 ...
## $ Received_TopDress : chr [1:528] "NO" "NO" "NO" "NO" ...
## $ tare           : num [1:528] 1220 1220 1220 1220 1220 1220 1220 1220 1220 1220 ...
## $ FW1PlusTare     : num [1:528] 4360 4818 5376 5598 4852 ...
## $ FW2PlusTare     : num [1:528] 3254 1220 1220 1220 1220 ...
## $ SSFWPlusTare    : num [1:528] 2324 1814 1994 2126 2106 ...
## $ SSODW          : num [1:528] 632 479 570 562 583 ...
## $ HarvestBagPlusTie : num [1:528] 162 176 169 121 120 ...
## $ Grain1PlusPaperBag1 : num [1:528] 230 173 211 233 265 ...
## $ PaperBag1       : num [1:528] 7.96 7.96 7.96 7.96 7.96 7.96 7.96 7.96 7.96 7.96 ...
## $ Grain2PlusPaperBag2 : num [1:528] 7.01 7.01 7.01 7.01 7.01 7.01 7.01 7.01 7.01 7.01 ...
## $ PaperBag2       : num [1:528] 7.01 7.01 7.01 7.01 7.01 7.01 7.01 7.01 7.01 7.01 ...
## $ StrawSampleSize : num [1:528] 8.25 7.96 7.76 7.89 8.45 ...
## $ StrawN          : num [1:528] 63.5 36 46.1 55.9 39.7 ...
## $ GrainSampleSize : num [1:528] 4.97 5.52 5.23 5.72 5.82 ...
## $ GrainN          : num [1:528] 60.5 50 56 66.7 51.1 ...
## $ DaysPI2Harvest   : num [1:528] 76 76 76 76 76 76 76 76 76 76 ...
## $ SeedTray1.1      : num [1:528] 243 243 243 243 243 243 243 243 243 243 ...
## $ SeedTray1.2      : num [1:528] 0.6 0.6 0.6 0.6 0.6 0.6 0.6 0.6 0.6 0.6 ...
## $ Grain3PlusSeedTray1 : num [1:528] 470 410 449 471 504 ...
## $ MoistureContentGrain3: num [1:528] 11.2 11.2 11.2 11.2 11.2 11.2 11.2 11.2 11.2 11.2 ...
## $ SeedTray2        : num [1:528] 244 244 244 244 244 ...
## $ Grain4PlusSeedTray2 : num [1:528] 254 254 254 254 254 ...

```

```
## $ SeasonalNRate_f      : Factor w/ 21 levels "0","25","30",...: 18 1 12 14 4 7 7 12 1 14 ...
## $ 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/ 4 levels "0","25","34",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ SeasonalNRate_kgha_f : Factor w/ 22 levels "0","25","34",...: 18 1 11 15 4 7 7 11 1 15 ...

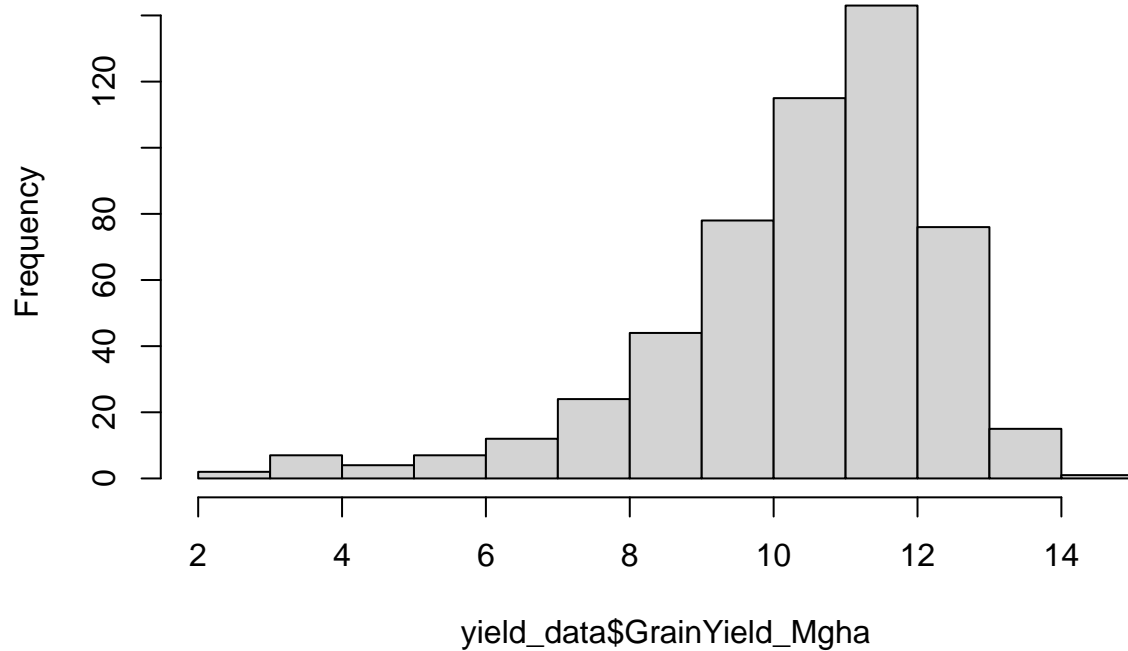
yield_data <- yield_data %>%
  mutate(
    FW1net = FW1PlusTare - tare,
    FW2net = FW2PlusTare - tare,
    TotalFW = FW1net + FW2net,
    SSFWnet = SSFWPlusTare - tare,
    Ratio = SSFWnet / TotalFW,
    SSODWnet = SSODW - HarvestBagPlusTie,
    SeedTray1 = SeedTray1.1 + SeedTray1.2, #adds the decimal 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)
```

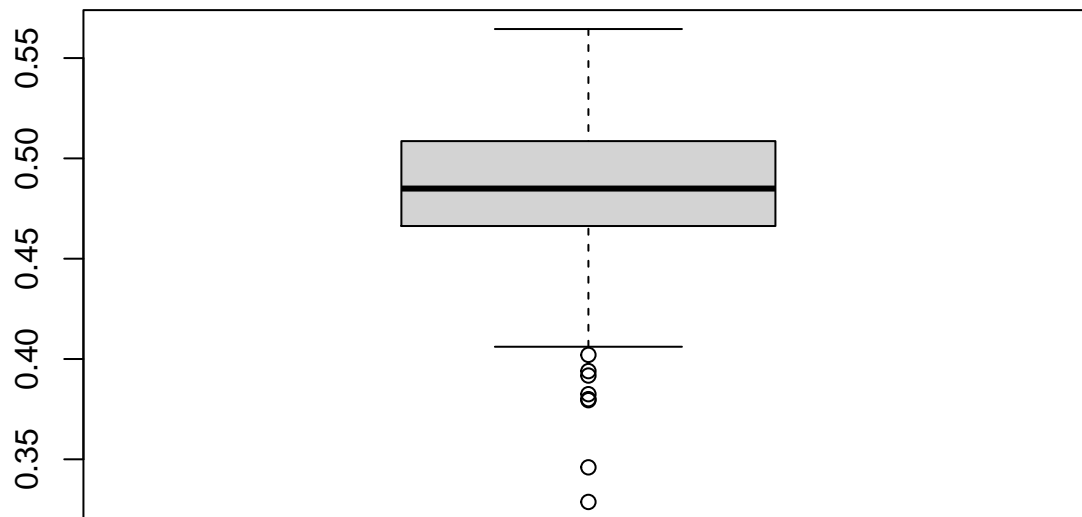



```
hist(yield_data$GrainYield_Mgha)
```

Histogram of yield_data\$GrainYield_Mgha

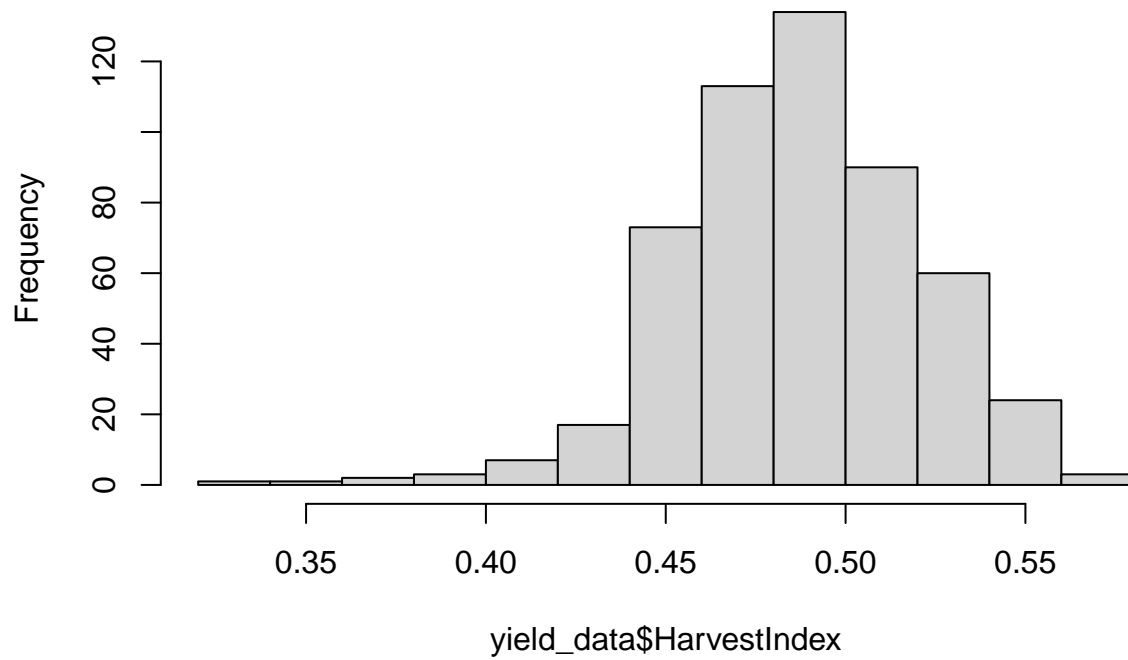


```
boxplot(yield_data$HarvestIndex)
```

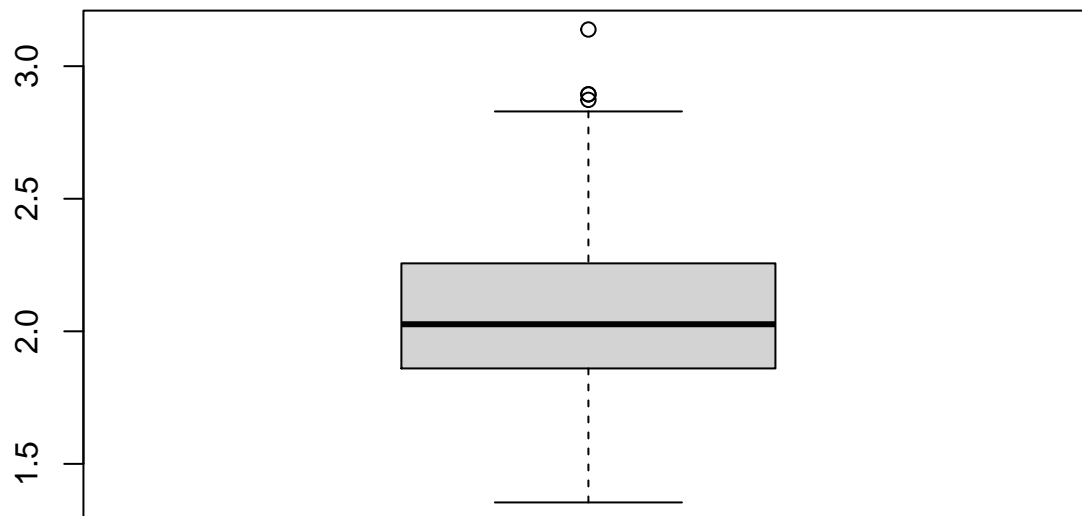


```
hist(yield_data$HarvestIndex)
```

Histogram of yield_data\$HarvestIndex

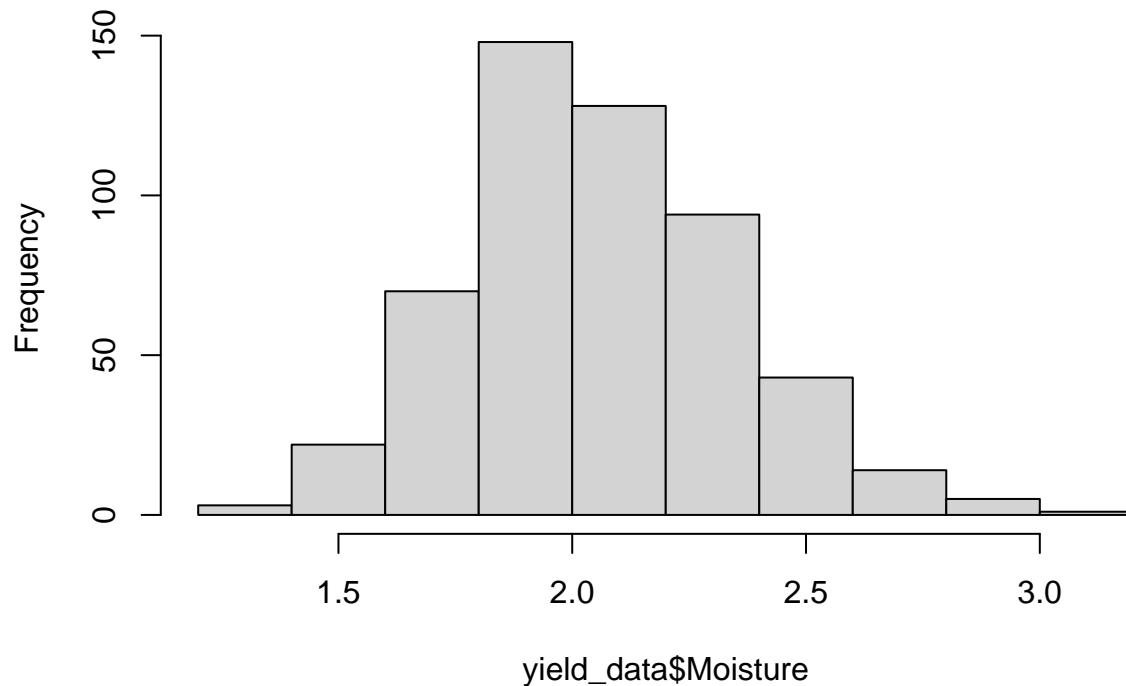


```
boxplot(yield_data$Moisture)
```



```
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

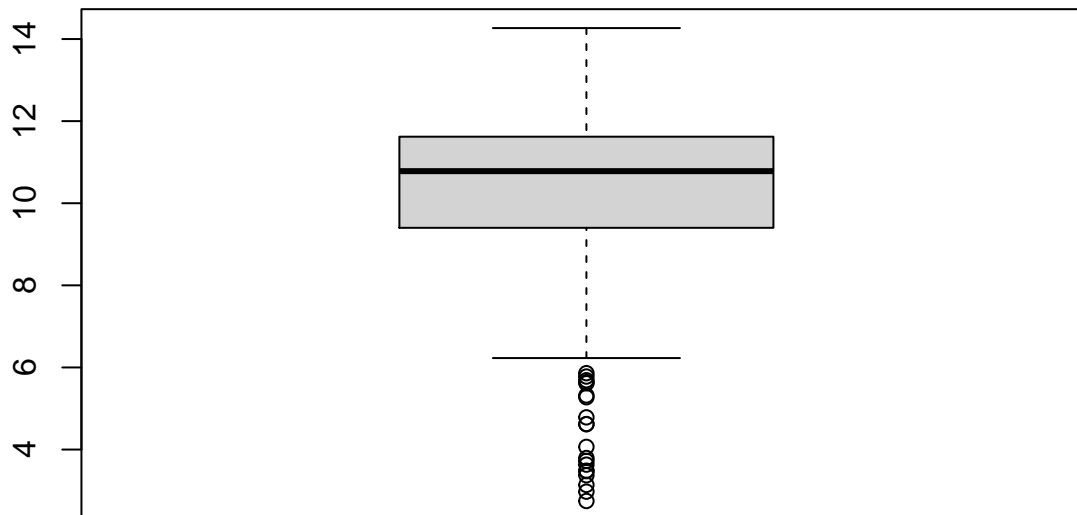
gs_data$site_year <- as.factor(gs_data$site_year)

str(gs_data , give.attr = FALSE)

## rowwise_df[,13] [528 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 1 1 2 2 2 3 3 3 4 ...
## $ Block          : Factor w/ 4 levels "1","2","3","4": 1 1 1 1 1 1 1 1 1 1 ...
## $ MainPlot       : Factor w/ 7 levels "1","2","3","4",...: 1 1 1 2 2 2 3 3 3 4 ...
## $ SubPlot        : Factor w/ 3 levels "1","2","3": 2 1 3 3 2 1 1 3 2 1 ...
## $ N_level_kgha    : num [1:528] 225 225 225 0 0 0 125 125 125 175 ...
## $ N_level_kgha_f  : Factor w/ 11 levels "0","45","75",...: 10 10 10 1 1 1 5 5 5 8 ...
## $ TopDress_kgha   : num [1:528] 0 25 50 0 25 50 0 25 50 0 ...
## $ TopDress_kgha_f: Factor w/ 4 levels "0","25","34",...: 1 2 4 1 2 4 1 2 4 1 ...
## $ PI_N_Uptake     : num [1:528] 158 158 158 78 78 ...
## $ NDVI            : num [1:528] 0.78 0.78 0.78 0.49 0.49 0.49 0.68 0.68 0.68 0.79 ...
## $ GrainYield_Mgha: num [1:528] 11.4 12.2 12.1 11 10.9 ...

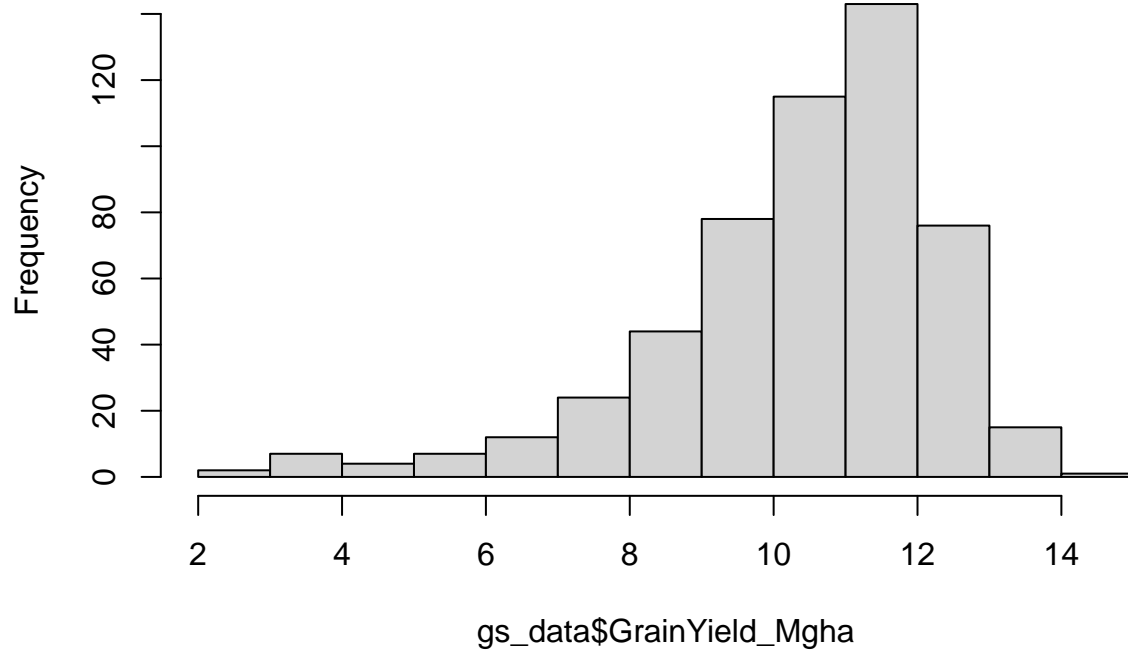
boxplot(gs_data$GrainYield_Mgha)

```



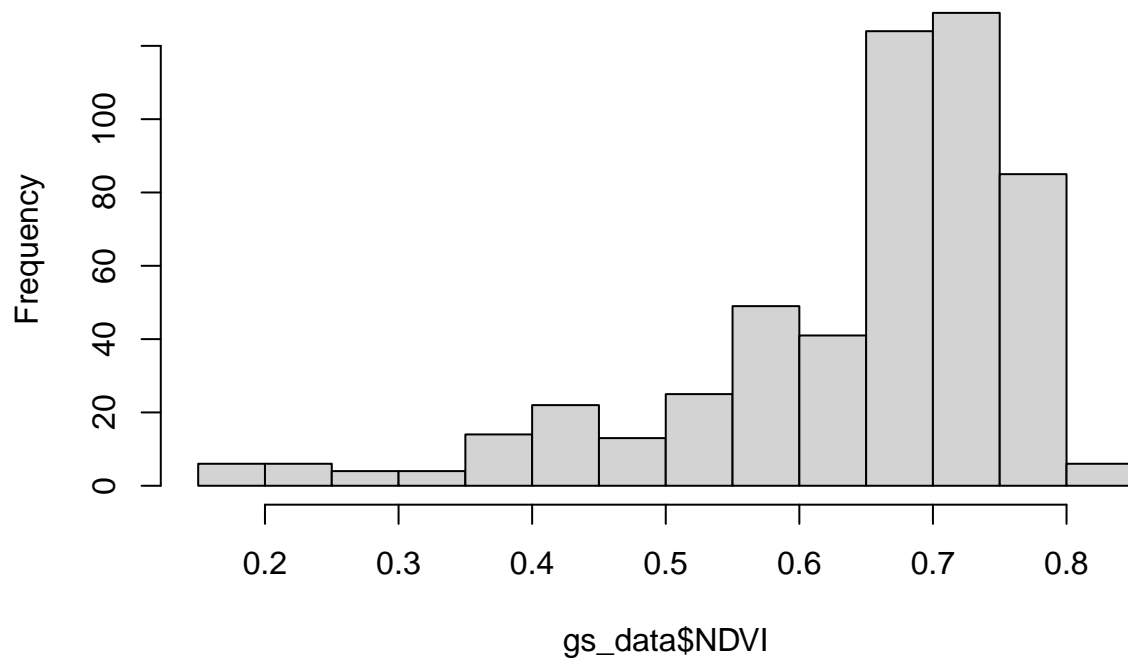
```
hist(gs_data$GrainYield_Mgha)
```

Histogram of gs_data\$GrainYield_Mgha

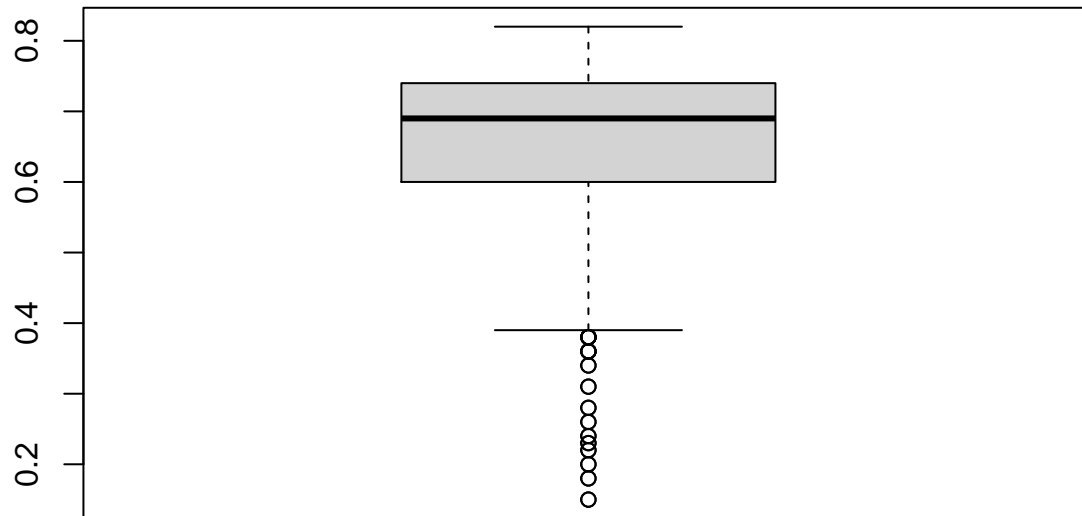


```
hist(gs_data$NDVI)
```

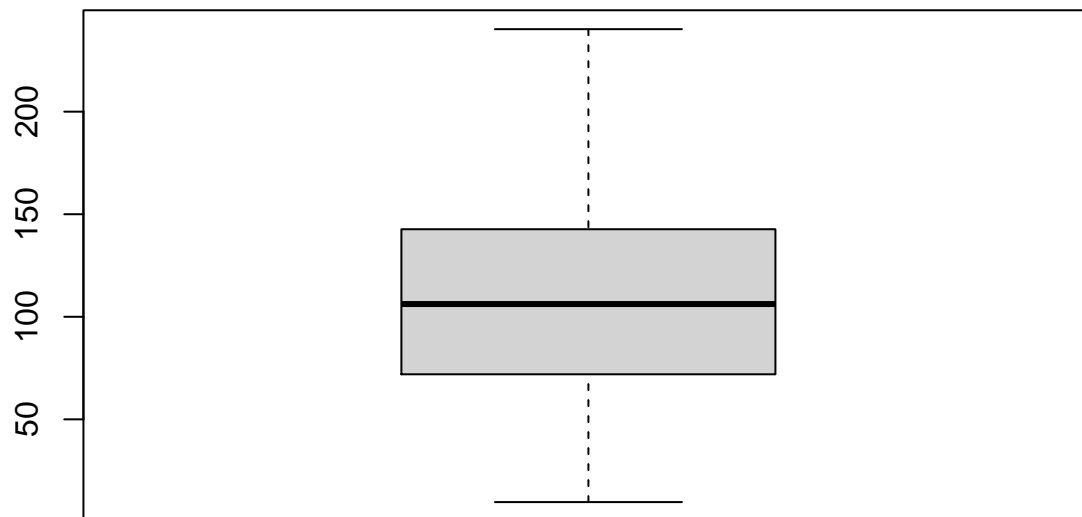
Histogram of gs_data\$NDVI



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

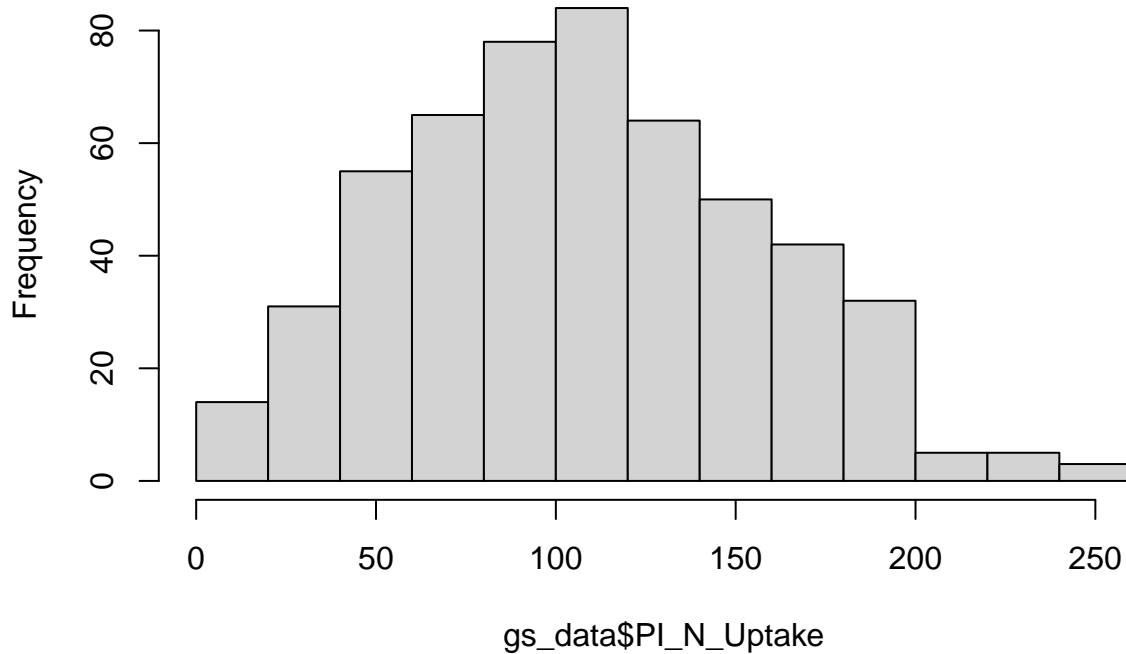


```
boxplot(gs_data$PI_N_Uptake)
```



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

Histogram of gs_data\$PI_N_Uptake



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

##Data for Table 3

```
table_3_data <- gs_data %>%
  select(site_year , TopDress_kgha_f , GrainYield_Mgha) %>%
  group_by(site_year , TopDress_kgha_f) %>%
  summarise(GrainYield_min = min(GrainYield_Mgha),
            GrainYield_max = max(GrainYield_Mgha),
            GrainYield_mean = mean(GrainYield_Mgha)) %>%
  ungroup()
```

`summarise()` has grouped output by 'site_year'. You can override using the `.groups` argument.

```
table_3_data$GrainYield_min <- round(table_3_data$GrainYield_min , digits = 1)
table_3_data$GrainYield_max <- round(table_3_data$GrainYield_max , digits = 1)
table_3_data$GrainYield_mean <- round(table_3_data$GrainYield_mean , digits = 1)
```

Calculating GS RI

```
max_gs_ndvi <- gs_data %>%
  filter(N_level_kgha_f %in% c(225, 235) & TopDress_kgha_f == 0) %>%
  select(site_year , NDVI) %>%
  group_by(site_year) %>%
  summarise(mean_NDVI = mean(NDVI))

max_gs_ndvi$mean_NDVI <- round(max_gs_ndvi$mean_NDVI , digits = 2)

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

```

nic17maxNDVI <- as.numeric(nic17maxNDVI)
nic17maxNDVI

## [1] 0.78

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

## [1] 0.79

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

## [1] 0.74

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

## [1] 0.78

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

## [1] 0.73

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

## [1] 0.76

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] 0.61

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

## [1] 0.75

```



```

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

```

```
## [1] 0.67
```

```

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)

```

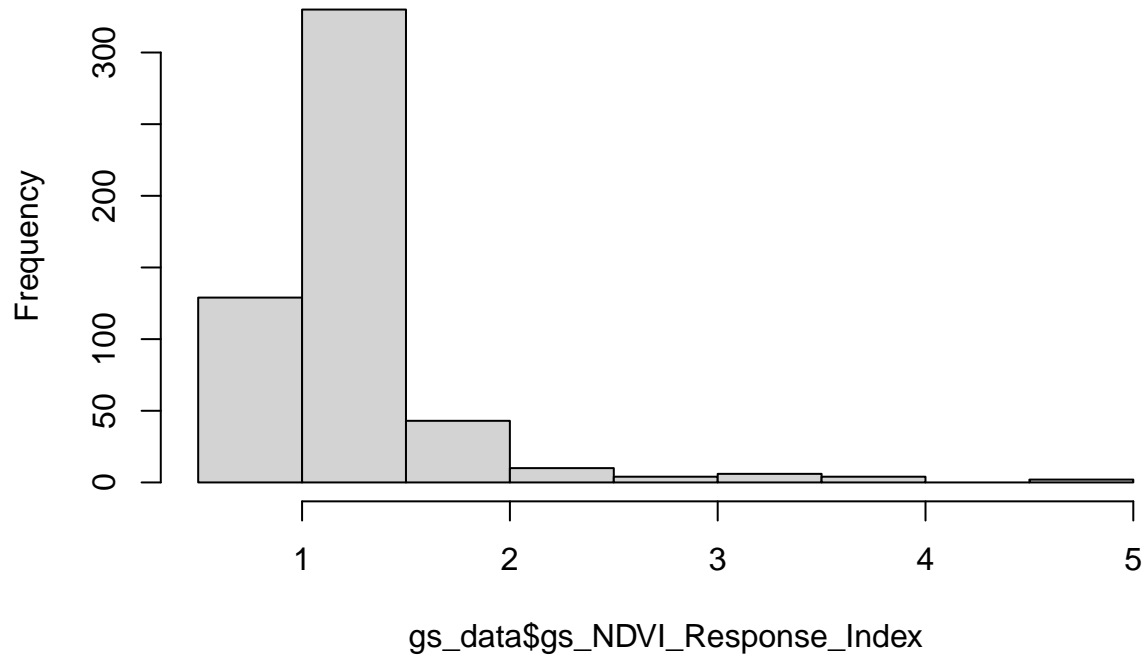
```

## rowwise_df[,15] [528 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 1 1 2 2 2 3 3 3 4 ...
## $ Block          : Factor w/ 4 levels "1","2","3","4": 1 1 1 1 1 1 1 1 1 1 ...
## $ MainPlot        : Factor w/ 7 levels "1","2","3","4",...: 1 1 1 2 2 2 3 3 3 4 ...
## $ SubPlot         : Factor w/ 3 levels "1","2","3": 2 1 3 3 2 1 1 3 2 1 ...
## $ N_level_kgha     : num [1:528] 225 225 225 0 0 0 125 125 125 175 ...
## $ N_level_kgha_f   : Factor w/ 11 levels "0","45","75",...: 10 10 10 1 1 1 5 5 5 8 ...
## $ TopDress_kgha    : num [1:528] 0 25 50 0 25 50 0 25 50 0 ...
## $ TopDress_kgha_f  : Factor w/ 4 levels "0","25","34",...: 1 2 4 1 2 4 1 2 4 1 ...
## $ PI_N_Uptake      : num [1:528] 158 158 158 78 78 ...
## $ NDVI             : num [1:528] 0.78 0.78 0.78 0.49 0.49 0.49 0.68 0.68 0.68 0.79 ...
## $ GrainYield_Mgha  : num [1:528] 11.4 12.2 12.1 11 10.9 ...
## $ max_NDVI         : num [1:528] 0.78 0.78 0.78 0.78 0.78 0.78 0.78 0.78 0.78 0.78 ...
## $ gs_NDVI_Response_Index: num [1:528] 1 1 1 1.59 1.59 ...

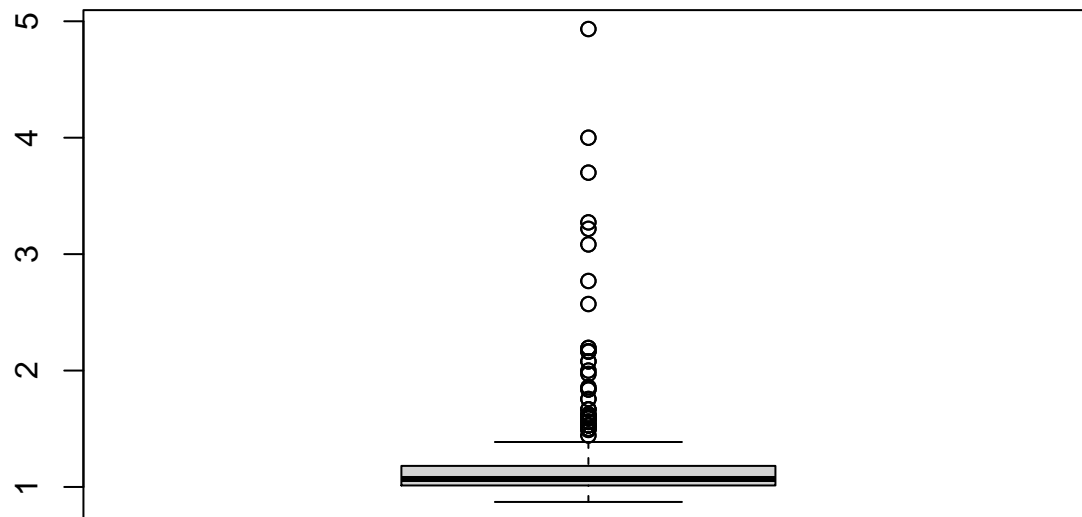
```

```
hist(gs_data$gs_NDVI_Response_Index)
```

Histogram of gs_data\$gs_NDVI_Response_Index

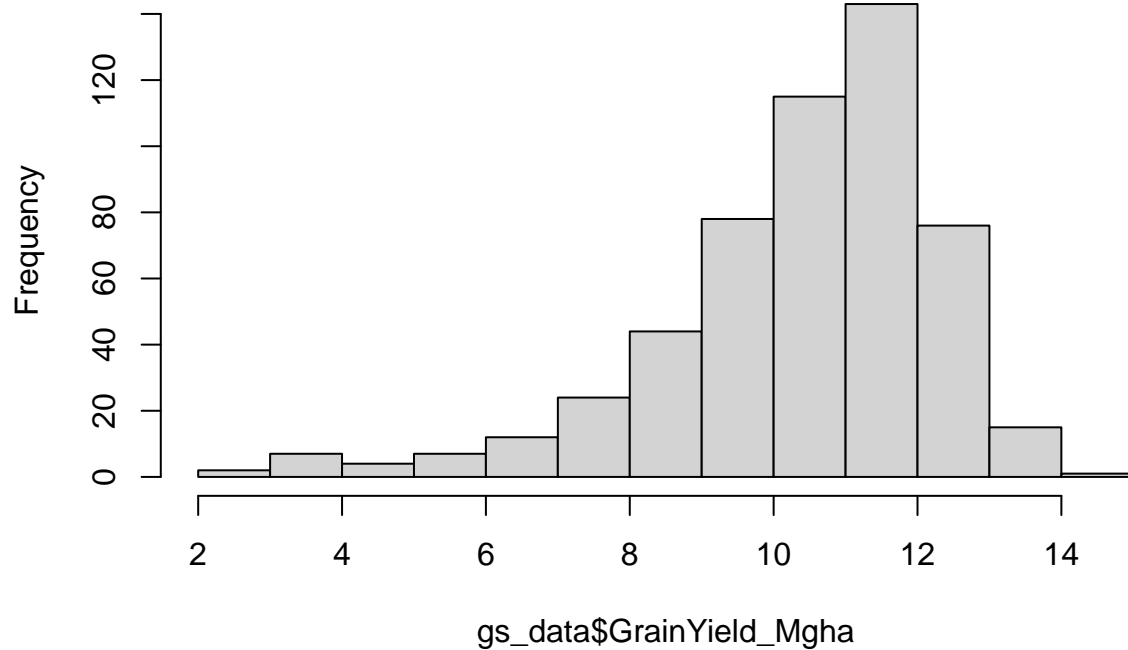


```
boxplot(gs_data$gs_NDVI_Response_Index)
```

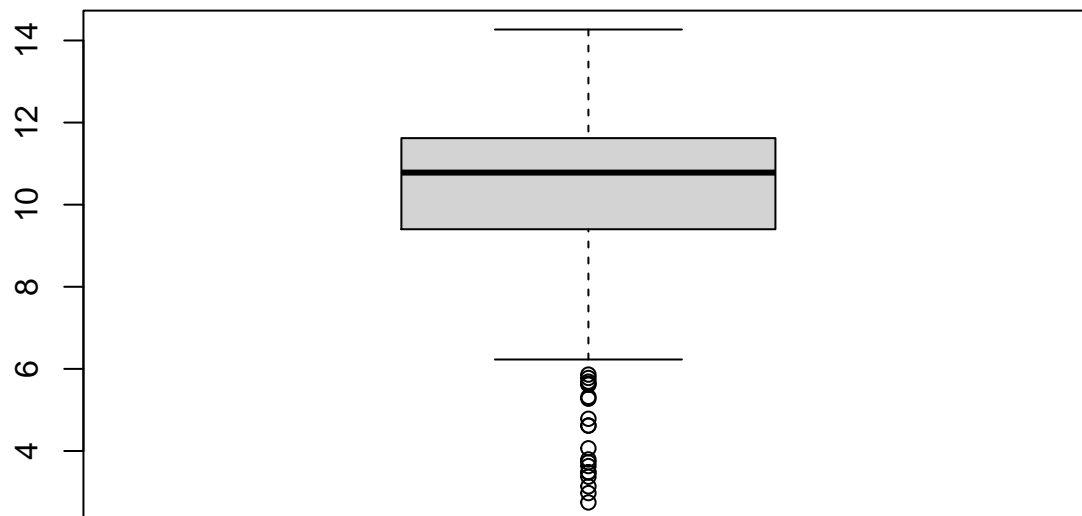


```
hist(gs_data$GrainYield_Mgha)
```

Histogram of gs_data\$GrainYield_Mgha



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



```
gs_data$gs_NDVI_Response_Index[gs_data$gs_NDVI_Response_Index < 1] <- 1 #converts values less than 1, t
```

```
##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.

drone_data <- drone_data %>%
  filter(N_level_kgha != 275)

str(drone_data , give.attr = FALSE)

## spec_tbl_df[,38] [240 x 38] (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 ...
## $ MainPlot       : 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 ...
## $ 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 0.0598 ...
## $ 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 0.0293 ...
## $ 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 0.0873 ...
## $ edgedmedian     : 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 ...

drone_data <- drone_data %>%
  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)

## spec_tbl_df[,39] [240 x 39] (S3: spec_tbl_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 ...
## $ 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
## $ edgedmedian     : 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

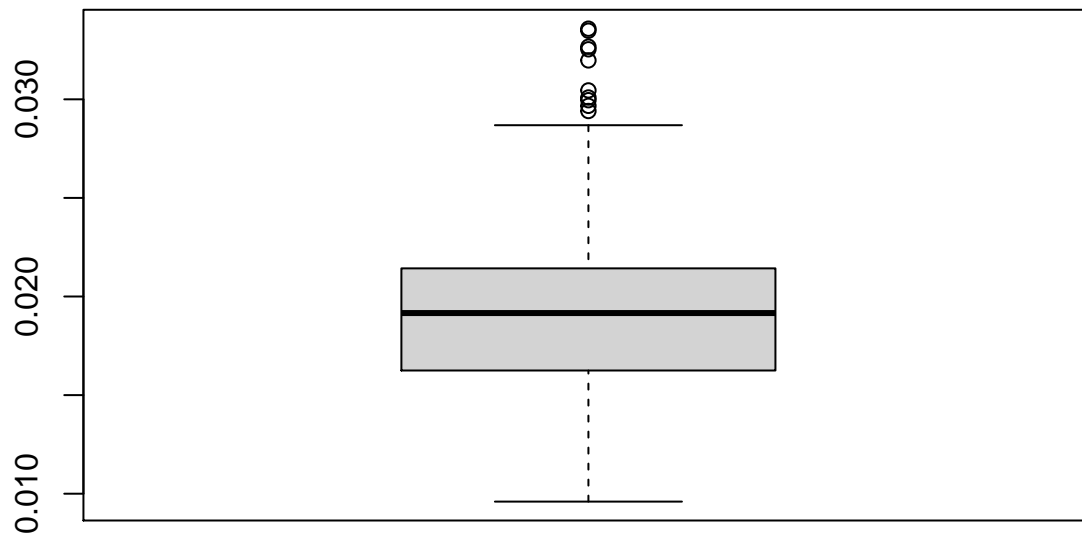
str(drone_data , give.attr = FALSE)

## tibble[,19] [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.
## $ 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.
## $ nirmean           : num [1:240] 0.55 0.357 0.482 0.537 0.431 ...

#visualize drone_data to look for outliers

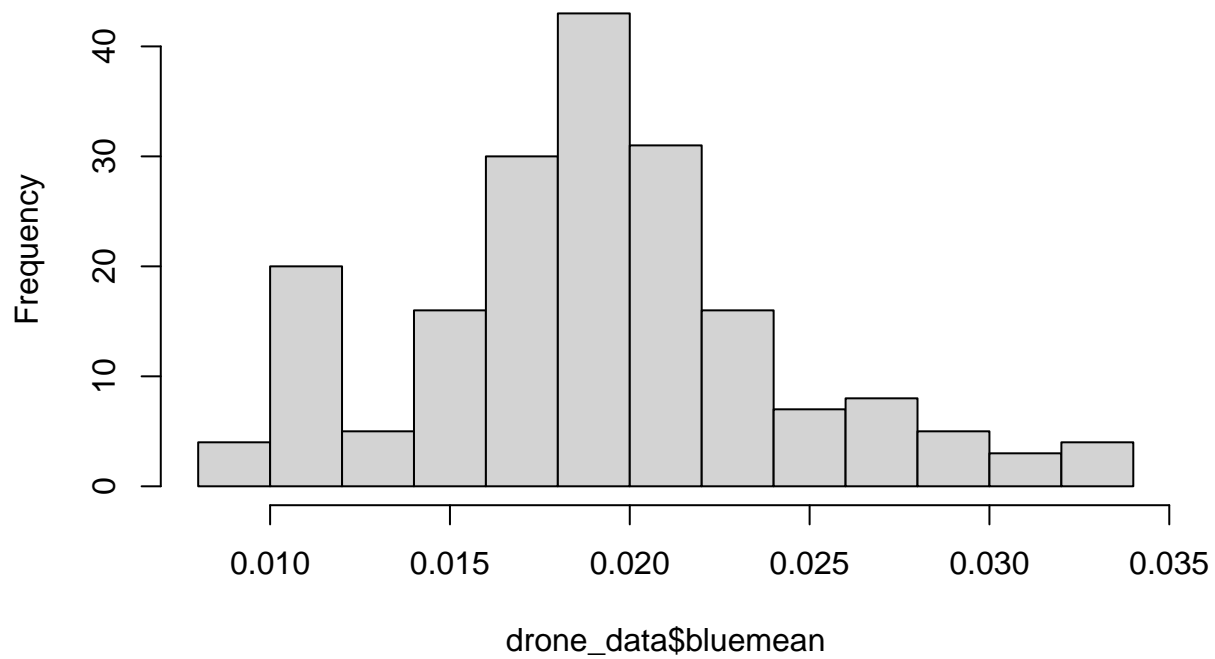
boxplot(drone_data$bluemean)

```

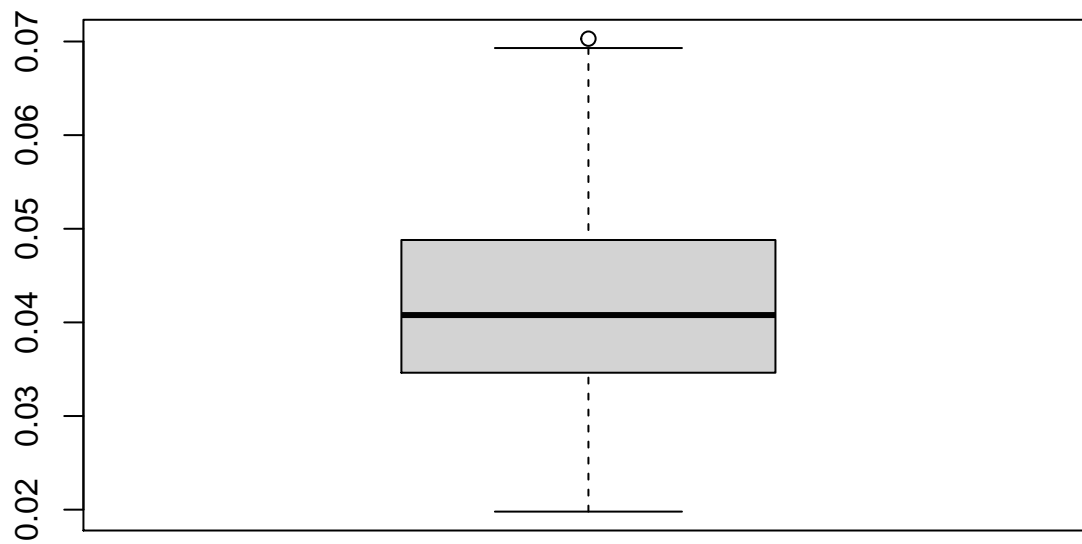


```
hist(drone_data$bluemean)
```

Histogram of drone_data\$bluemean

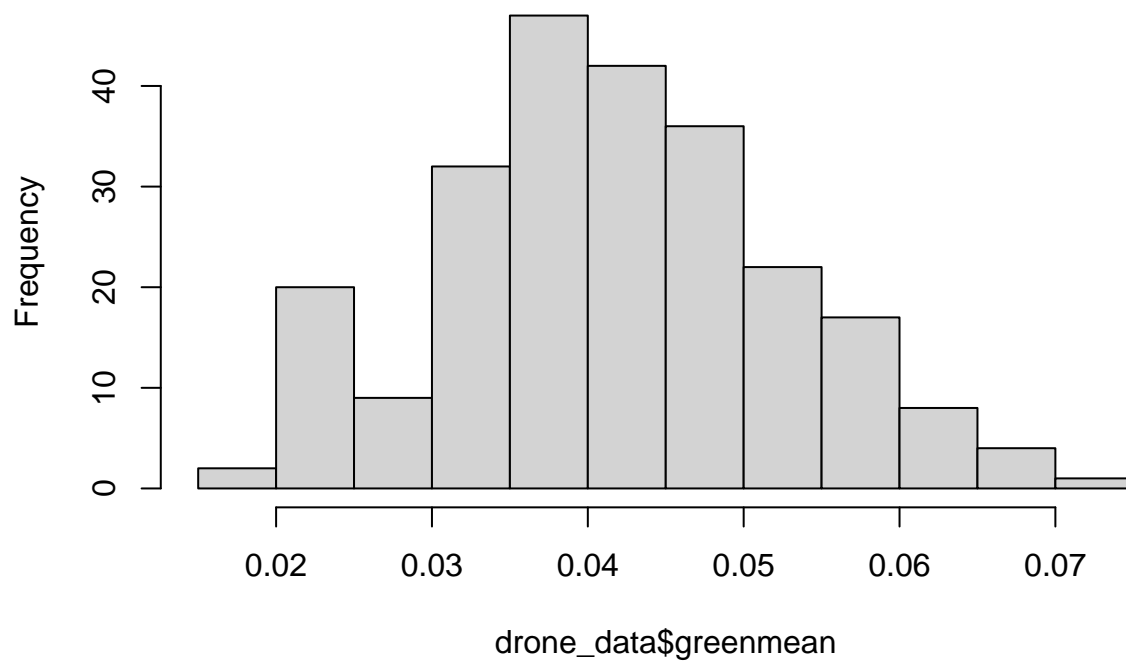


```
boxplot(drone_data$greenmean)
```

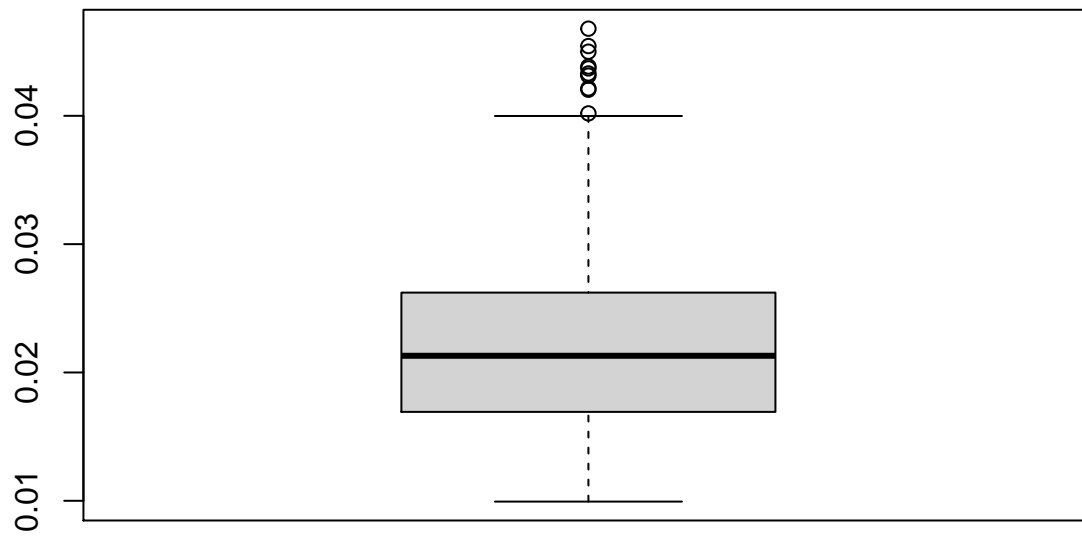


```
hist(drone_data$greenmean)
```

Histogram of drone_data\$greenmean

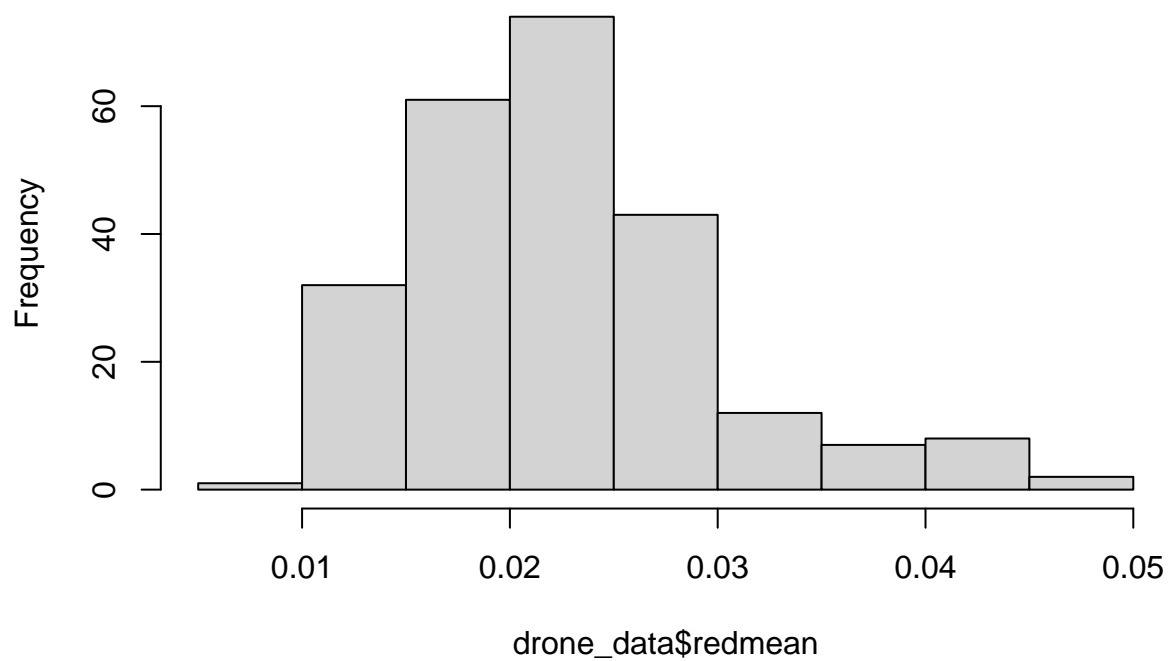


```
boxplot(drone_data$redmean)
```

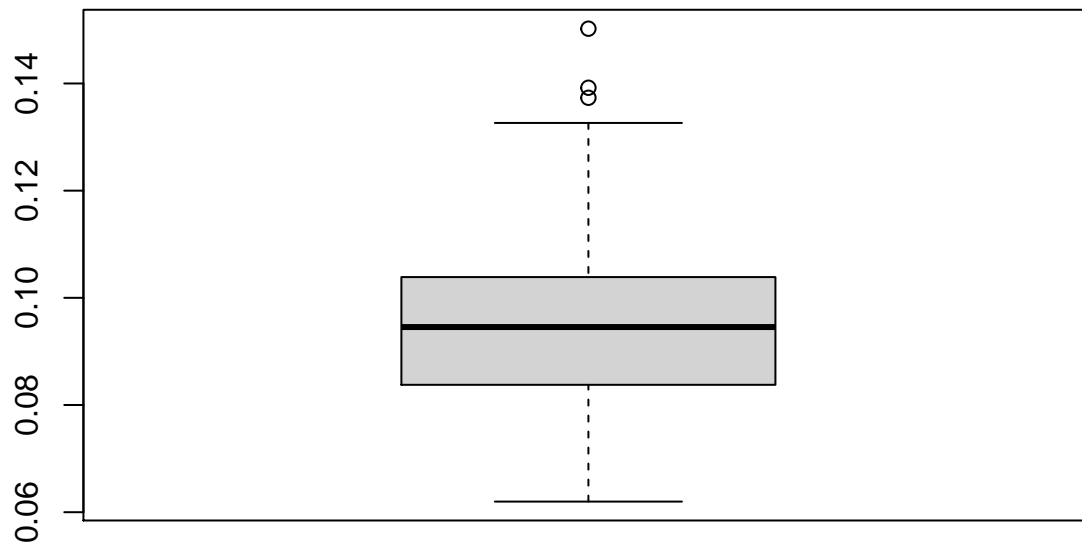



```
hist(drone_data$redmean)
```

Histogram of drone_data\$redmean

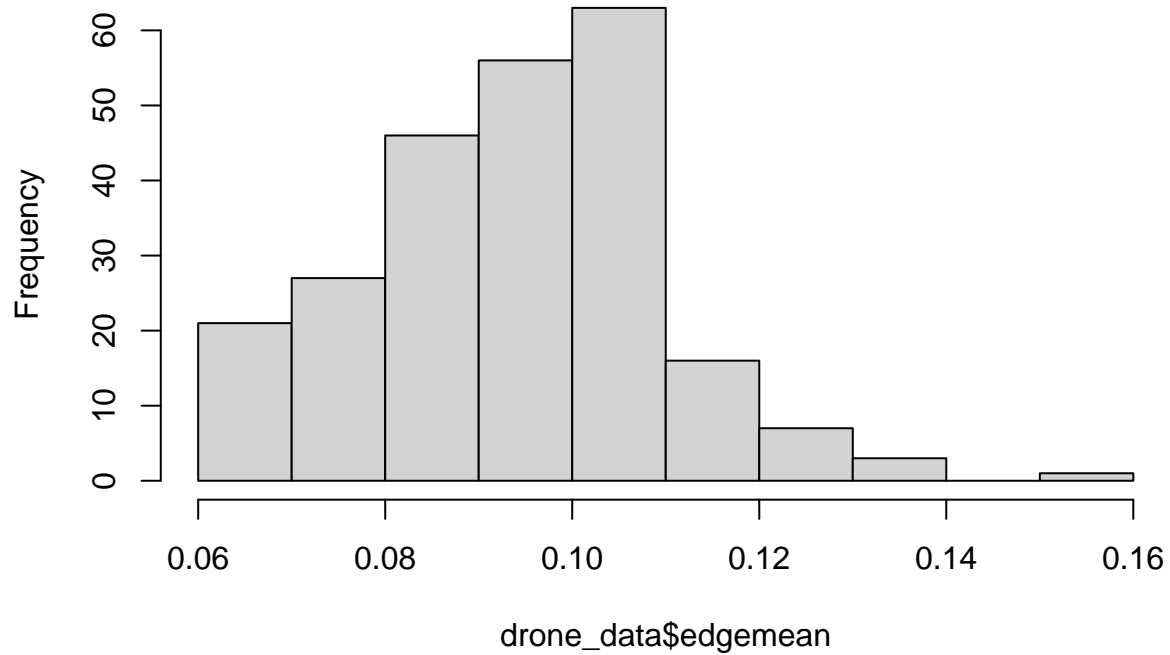


```
boxplot(drone_data$edgemean)
```

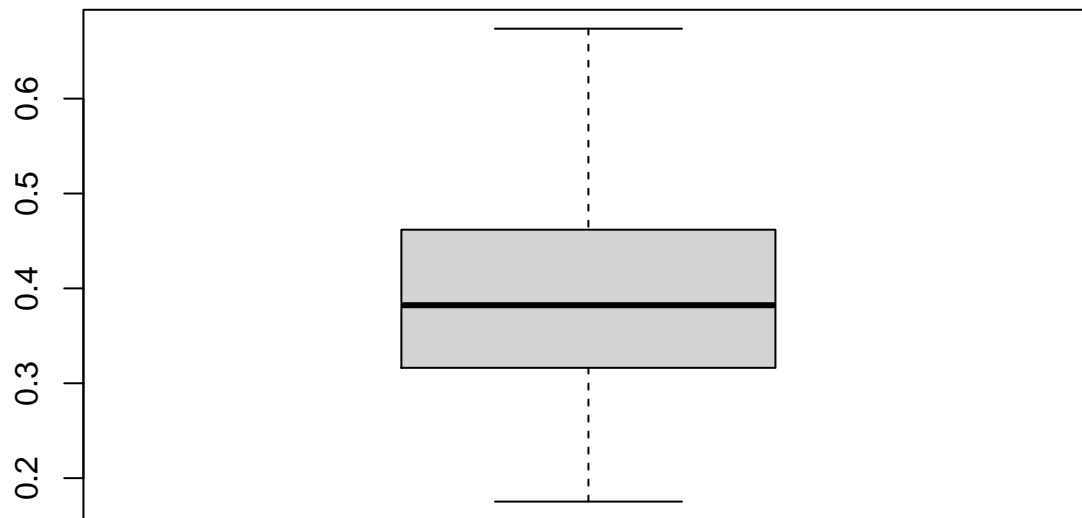


```
hist(drone_data$edgemean)
```

Histogram of drone_data\$edgemean

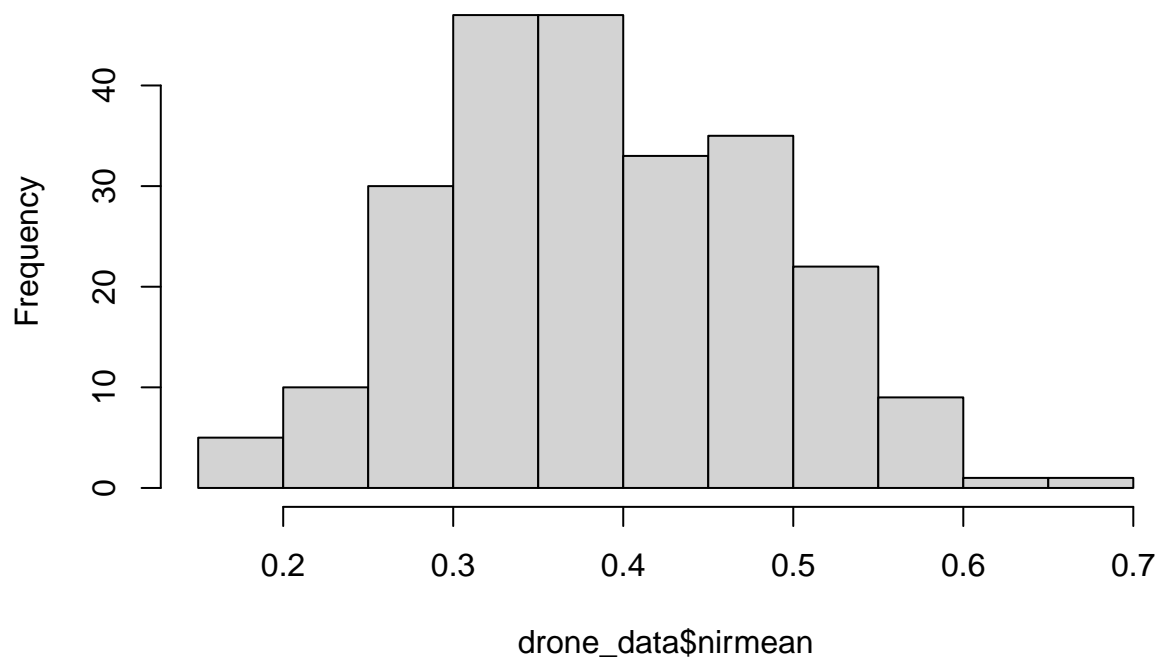


```
boxplot(drone_data$nirmean)
```



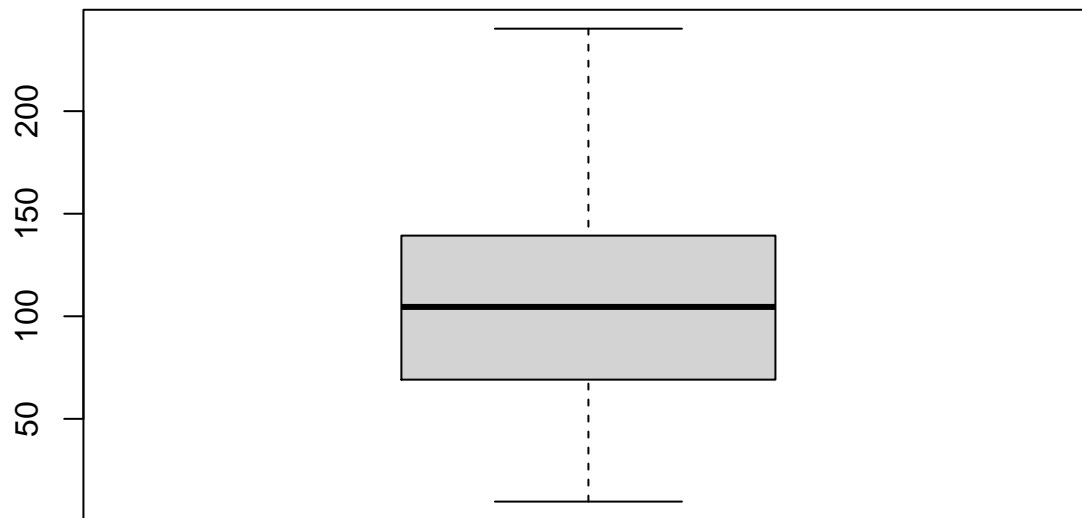
```
hist(drone_data$nirmean)
```

Histogram of drone_data\$nirmean



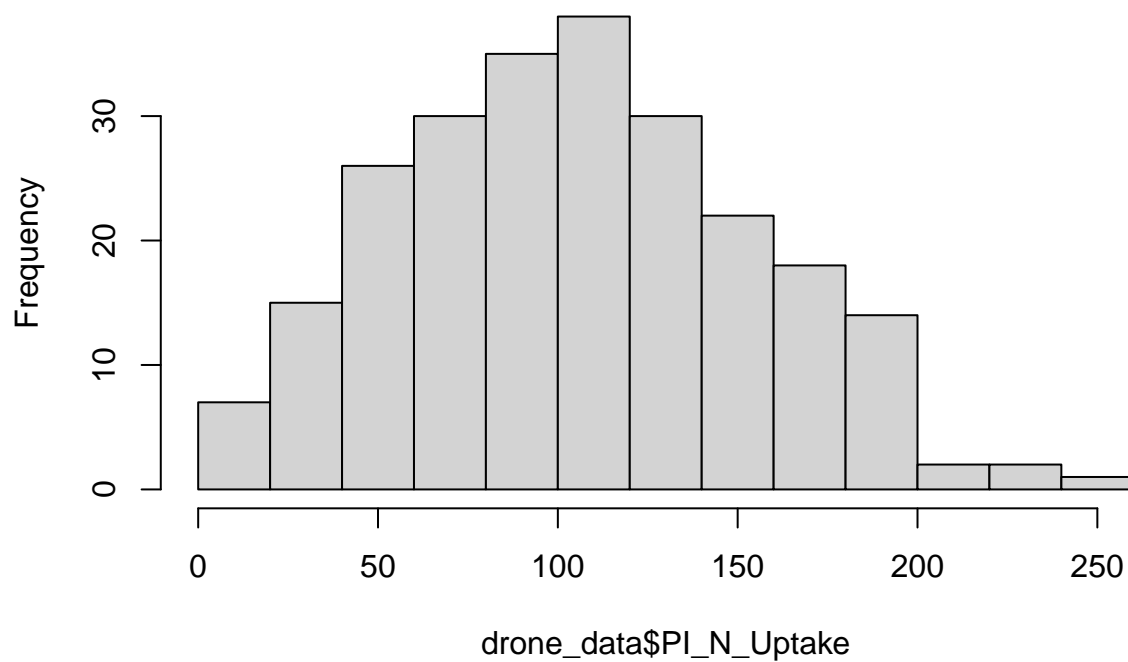
```
drone_data <- drone_data %>%
  mutate( biomass_dry_wt = biomass_plus_bag_g - (paper_bag_g * num_of_paper_bags) ,
           aboveground_biomass = (biomass_dry_wt / ring_size) * 10 , #ring size 0.5 m^2 biomass in kg per ha
           n_content = sample_N Ug / sample_weight_mg ,
           PI_N_Uptake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

boxplot(drone_data$PI_N_Uptake)
```



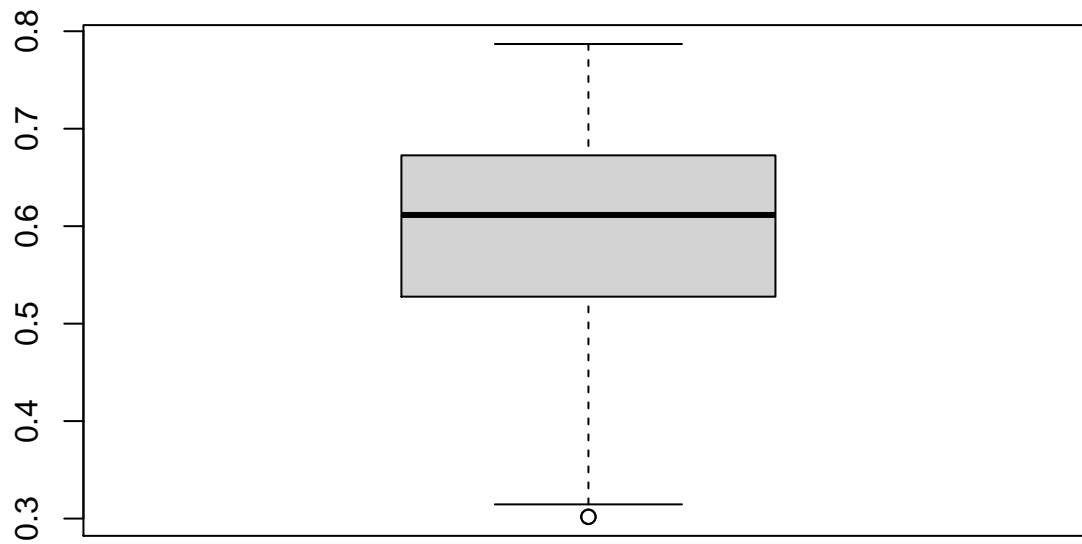
```
hist(drone_data$PI_N_Uptake)
```

Histogram of drone_data\$PI_N_Uptake



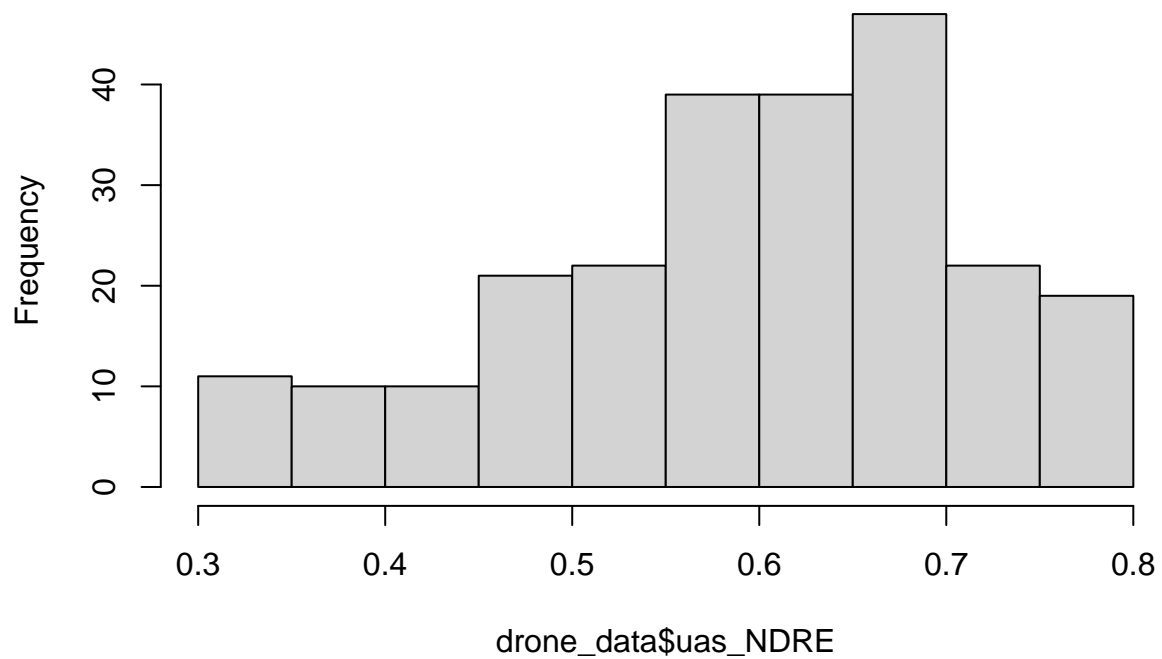
```
drone_data <- drone_data %>%
  mutate(uas_NDRE = ((nirmean - edgemean) / (nirmean + edgemean)) ,
         uas_NDVI = ((nirmean - redmean) / (nirmean + redmean))
         ) #calculates NDRE and NDVI

boxplot(drone_data$uas_NDRE)
```

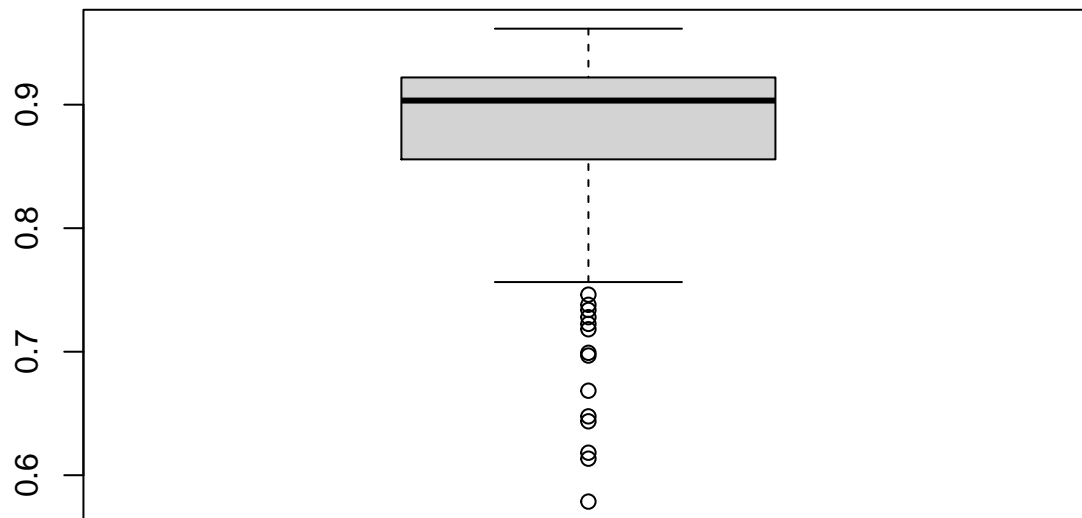


```
hist(drone_data$uas_NDRE)
```

Histogram of drone_data\$uas_NDRE

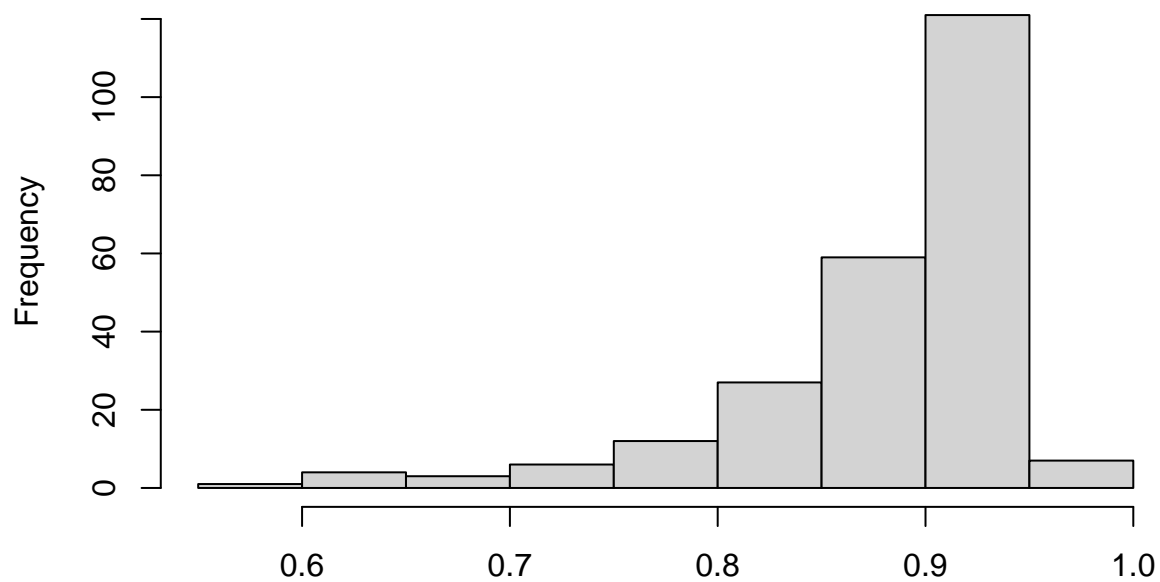


```
boxplot(drone_data$uas_NDVI)
```



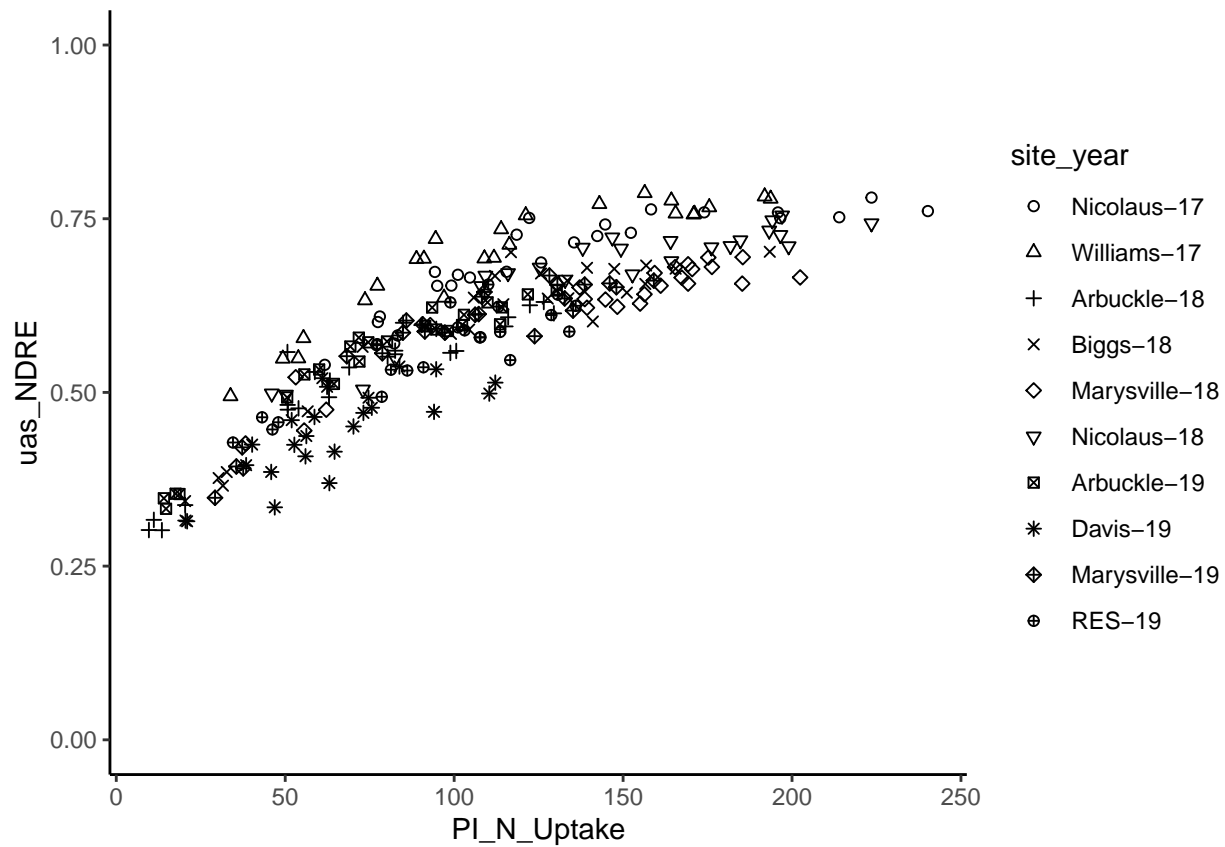
```
hist(drone_data$uas_NDVI)
```

Histogram of drone_data\$uas_NDVI

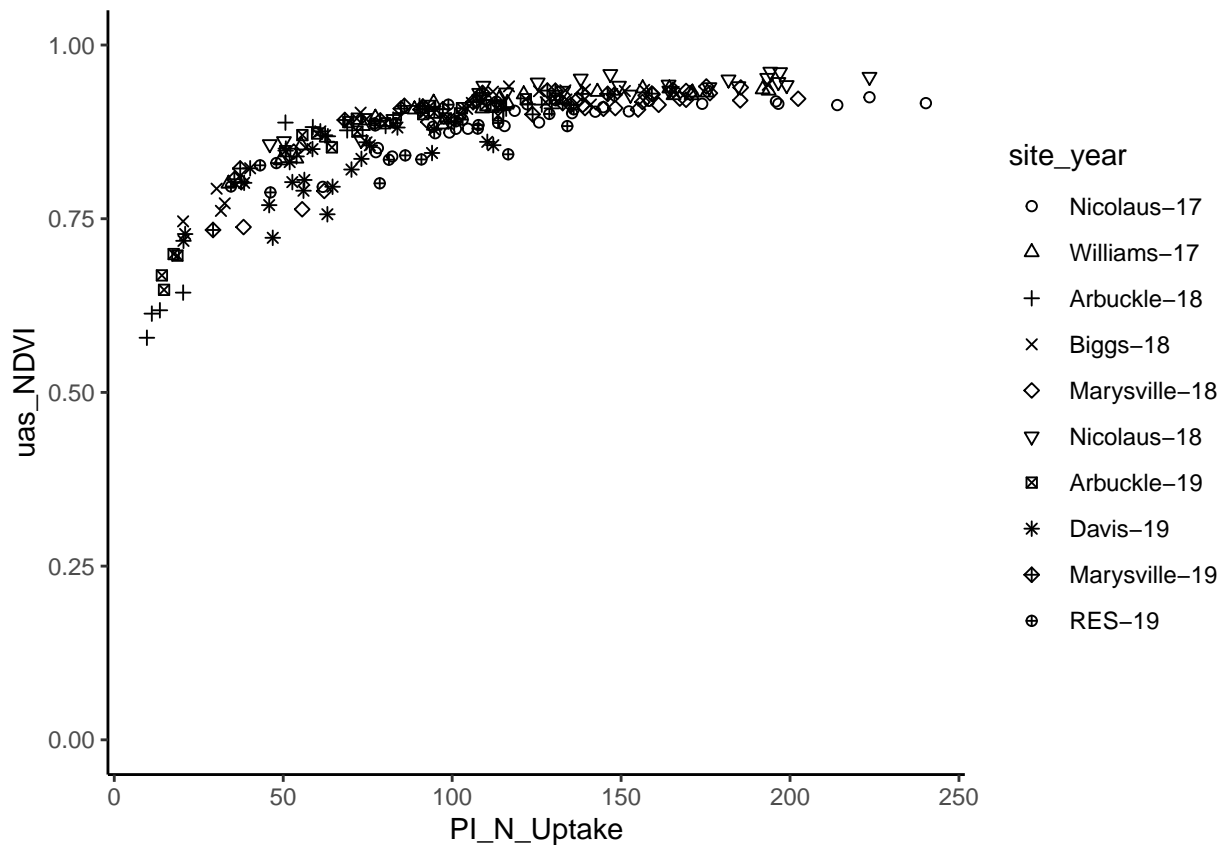


drone_data\$uas_NDVI

```
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))
```



```
drone_data$uas_NDRE <- round(drone_data$uas_NDRE , digits = 2) #rounds to the precision of the instrument
drone_data$uas_NDVI <- round(drone_data$uas_NDVI , digits = 2) #rounds to the precision of the instrument
```

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)) %>%
  select(site_year , uas_NDVI , uas_NDRE) %>%
  group_by(site_year) %>%
  summarise_all(.funs = mean) %>%
  ungroup()

max_drone_data$uas_NDVI <- round(max_drone_data$uas_NDVI , digits = 2)
max_drone_data$uas_NDRE <- round(max_drone_data$uas_NDRE , digits = 2)

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

```
## [1] 0.76
```

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

```
## [1] 0.92
```



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

```
## [1] 0.78
```

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

```
## [1] 0.94
```

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

```
## [1] 0.62
```

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

```
## [1] 0.91
```

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

```
## [1] 0.68
```

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

```
## [1] 0.93
```

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

```
## [1] 0.68
```

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

```
## [1] 0.93
```

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

```
## [1] 0.74
```

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

```
## [1] 0.96
```

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

```
## [1] 0.63
```

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

```
## [1] 0.91
```

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

```
## [1] 0.5
```

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

```
## [1] 0.86
```

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

```
## [1] 0.66
```

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

```
## [1] 0.93
```

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

```
## [1] 0.6
```

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

```
## [1] 0.88
```

```
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" ~ wil17maxuas_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 site
  )

drone_data <- drone_data %>%
  mutate(uas_NDVI_Response_Index = case_when(
    site_year == "Nicolaus-17" ~ nic17maxuas_NDVI / uas_NDVI,
    site_year == "Williams-17" ~ wil17maxuas_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

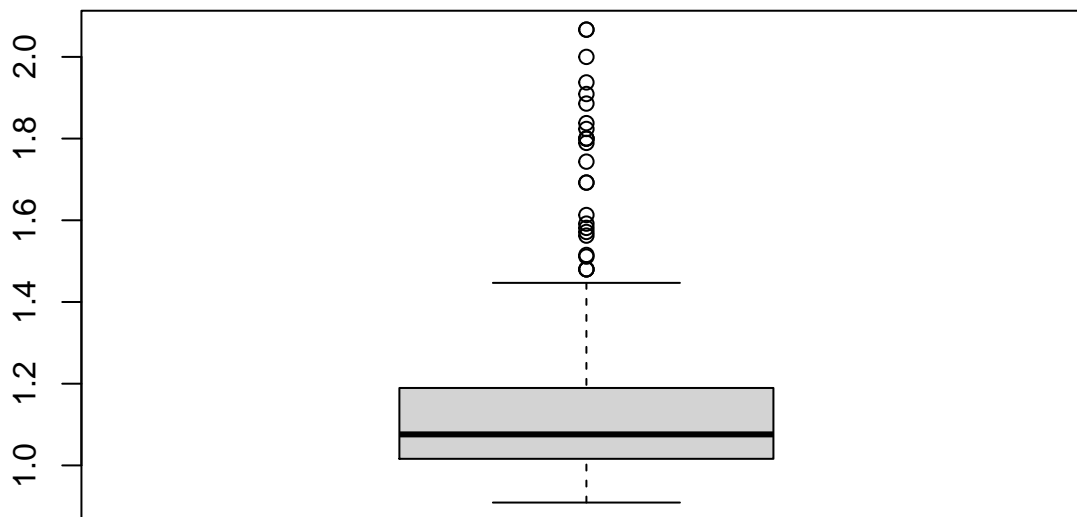
str(drone_data , give.attr = F)

## tibble[,29] [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 ...
## $ greenmean        : num [1:240] 0.0467 0.0581 0.0498 0.0488 0.0608 0.0589 0.0537 0.0488 0.05

```

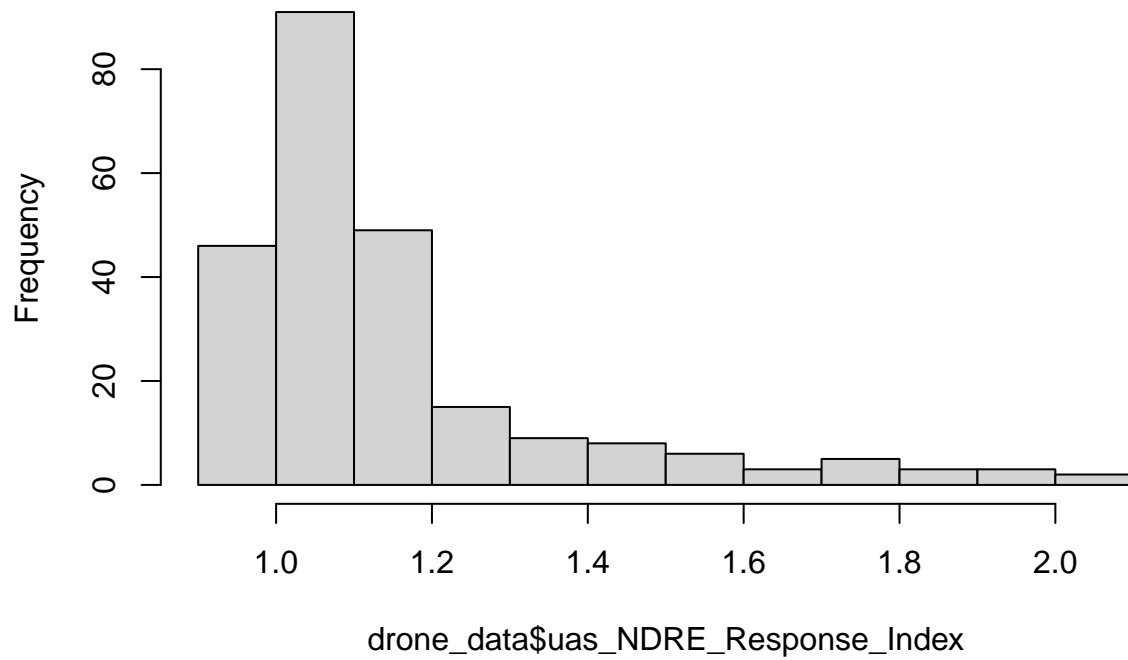
```
## $ redmean          : num [1:240] 0.023 0.0286 0.0243 0.0237 0.0292 0.0281 0.0263 0.0237 0.029
## $ edgemean         : num [1:240] 0.0738 0.0867 0.0768 0.0764 0.0903 0.0877 0.0812 0.0754 0.08
## $ nirmean          : num [1:240] 0.55 0.357 0.482 0.537 0.431 ...
## $ biomass_dry_wt    : num [1:240] 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] 0.76 0.61 0.73 0.75 0.65 0.67 0.67 0.73 0.6 0.76 ...
## $ uas_NDVI          : num [1:240] 0.92 0.85 0.9 0.92 0.87 0.88 0.88 0.91 0.85 0.92 ...
## $ max_uas_NDRE      : num [1:240] 0.76 0.76 0.76 0.76 0.76 0.76 0.76 0.76 0.76 0.76 ...
## $ uas_NDRE_Response_Index: num [1:240] 1 1.25 1.04 1.01 1.17 ...
## $ max_uas_NDVI      : num [1:240] 0.92 0.92 0.92 0.92 0.92 0.92 0.92 0.92 0.92 0.92 ...
## $ uas_NDVI_Response_Index: num [1:240] 1 1.08 1.02 1 1.06 ...
```

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

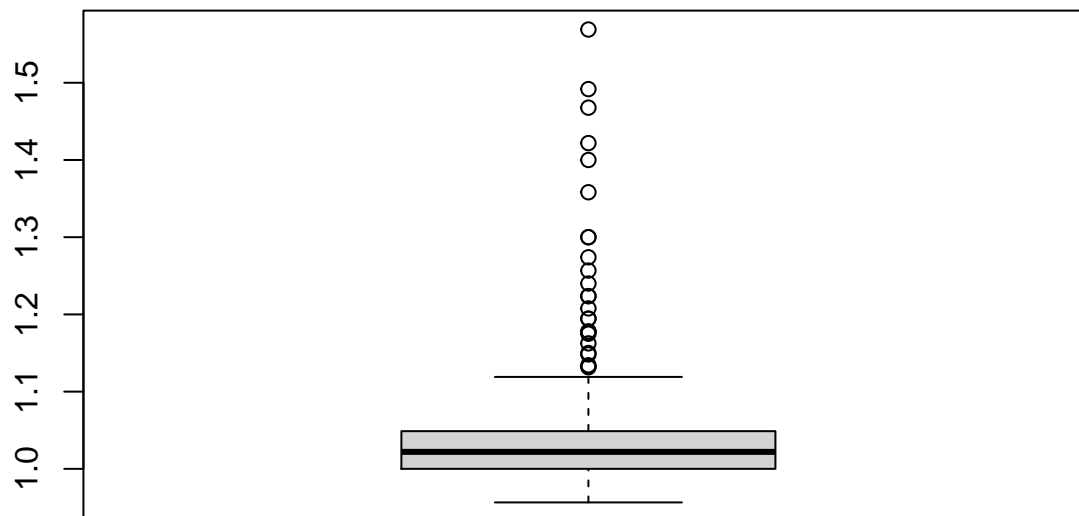


```
hist(drone_data$uas_NDRE_Response_Index)
```

Histogram of drone_data\$uas_NDRE_Response_Index

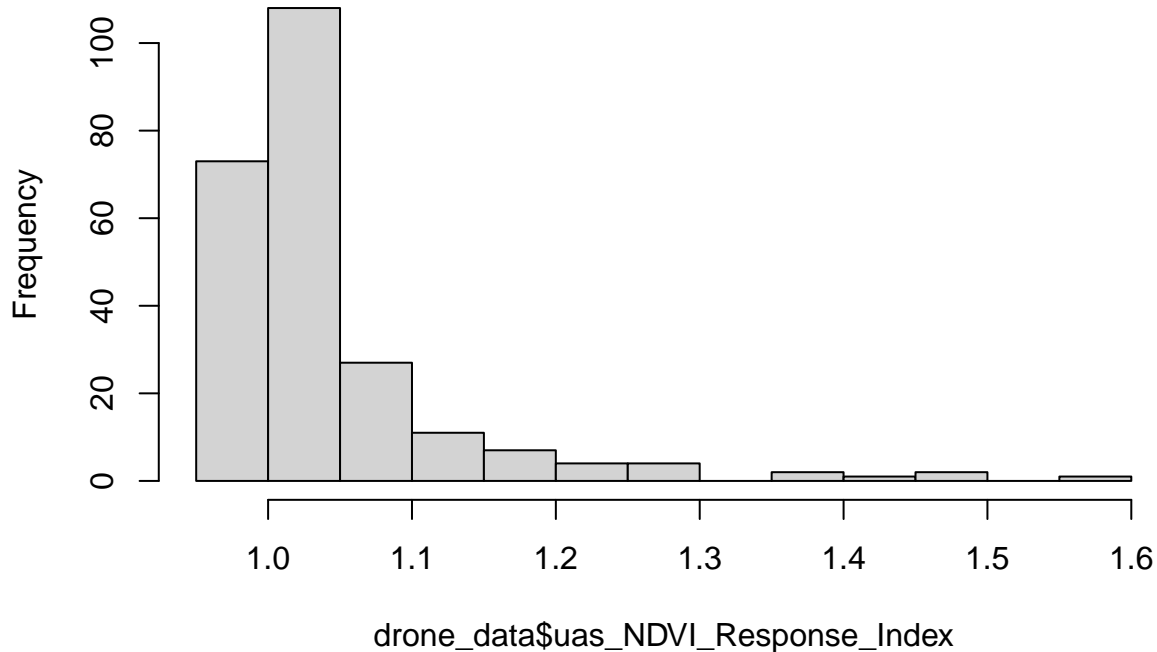


```
boxplot(drone_data$uas_NDVI_Response_Index)
```



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

Histogram of drone_data\$uas_NDVI_Response_Index



```
drone_data$uas_NDVI_Response_Index[drone_data$uas_NDVI_Response_Index < 1] <- 1 #converts values less than 1 to 1
drone_data$uas_NDRE_Response_Index[drone_data$uas_NDRE_Response_Index < 1] <- 1 #converts values less than 1 to 1
```

```
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[,12] [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] 0.76 0.61 0.73 0.75 0.65 0.67 0.67 0.73 0.6 0.76 ...
## $ uas_NDVI          : num [1:240] 0.92 0.85 0.9 0.92 0.87 0.88 0.88 0.91 0.85 0.92 ...
## $ uas_NDRE_Response_Index: num [1:240] 1 1.25 1.04 1.01 1.17 ...
## $ uas_NDVI_Response_Index: num [1:240] 1 1.08 1.02 1 1.06 ...
```

```
str(sUAS_yield_data , give.attr = F)
```

```
## tibble[,11] [528 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 ...
## $ 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 [1:528] 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 [1:528] 0 0 0 0 0 0 0 0 0 0 ...
## $ TopDress_kgha_f: Factor w/ 4 levels "0","25","34",...: 1 1 1 1 1 1 1 1 1 1 ...
## $ GrainYield_Mgha: num [1:528] 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[,16] [528 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 1 1 2 2 2 3 3 3 4 ...
## $ Block          : Factor w/ 4 levels "1","2","3","4": 1 1 1 1 1 1 1 1 1 1 ...
## $ MainPlot       : Factor w/ 7 levels "1","2","3","4",...: 1 1 1 2 2 2 3 3 3 4 ...
## $ SubPlot        : Factor w/ 3 levels "1","2","3": 2 1 3 3 2 1 1 3 2 1 ...
```

```
## $ N_level_kgha      : num [1:528] 225 225 225 0 0 0 125 125 125 175 ...
## $ N_level_kgha_f    : Factor w/ 11 levels "0","45","75",...: 10 10 10 1 1 1 5 5 5 8 ...
## $ TopDress_kgha     : num [1:528] 0 25 50 0 25 50 0 25 50 0 ...
## $ TopDress_kgha_f   : Factor w/ 4 levels "0","25","34",...: 1 2 4 1 2 4 1 2 4 1 ...
## $ PI_N_Uptake       : num [1:528] 158 158 158 78 78 ...
## $ uas_NDRE          : num [1:528] 0.76 0.76 0.76 0.61 0.61 0.61 0.73 0.73 0.73 0.75 ...
## $ uas_NDVI          : num [1:528] 0.92 0.92 0.92 0.85 0.85 0.85 0.9 0.9 0.9 0.92 ...
## $ GrainYield_Mgha   : num [1:528] 11.4 12.2 12.1 11 10.9 ...
## $ uas_NDRE_Response_Index: num [1:528] 1 1 1 1.25 1.25 ...
## $ uas_NDVI_Response_Index: num [1:528] 1 1 1 1.08 1.08 ...
```

##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")
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" ,
         SI_sq = (gs_NDVI_SI * gs_NDVI_SI)) %>%
  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" ,
         SI_sq = (uas_NDVI_SI * uas_NDVI_SI)) %>%
  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" ,
       SI_sq = (uas_NDRE_SI * uas_NDRE_SI)) %>%
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 ,
               SI_sq ,
               GrainYield_Mgha,
               PI_N_Uptake)

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

str(paper3_data , give.attr = F)

## rowwise_df[,17] [1,584 x 17] (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 1 1 2 2 2 3 3 3 4 ...
## $ Block          : Factor w/ 4 levels "1","2","3","4": 1 1 1 1 1 1 1 1 1 1 ...
## $ MainPlot       : Factor w/ 7 levels "1","2","3","4",...: 1 1 1 2 2 2 3 3 3 4 ...

```

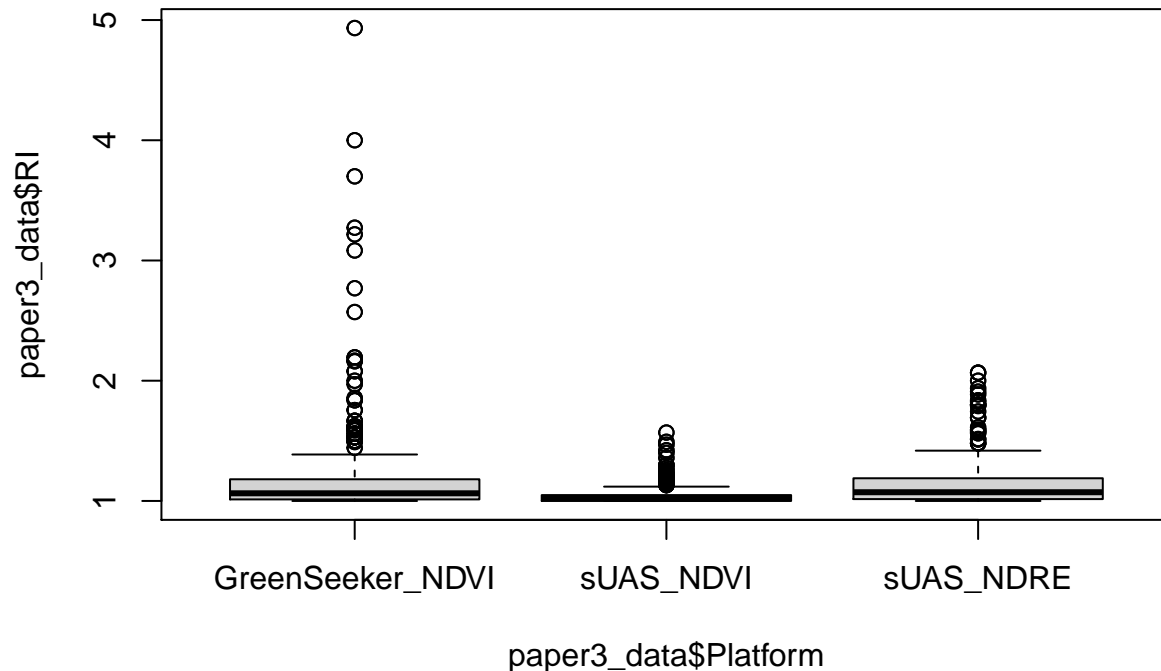
```
## $ SubPlot      : Factor w/ 3 levels "1","2","3": 2 1 3 3 2 1 1 3 2 1 ...
## $ N_level_kgha : num [1:1584] 225 225 225 0 0 0 125 125 125 175 ...
## $ N_level_kgha_f : Factor w/ 11 levels "0","45","75",...: 10 10 10 1 1 1 5 5 5 8 ...
## $ TopDress_kgha : num [1:1584] 0 25 50 0 25 50 0 25 50 0 ...
## $ TopDress_kgha_f: Factor w/ 4 levels "0","25","34",...: 1 2 4 1 2 4 1 2 4 1 ...
## $ Index        : num [1:1584] 0.78 0.78 0.78 0.49 0.49 0.49 0.68 0.68 0.68 0.79 ...
## $ RI           : num [1:1584] 1 1 1 1.59 1.59 ...
## $ SI           : num [1:1584] 1 1 1 0.628 0.628 ...
## $ SI_sq        : num [1:1584] 1 1 1 0.395 0.395 ...
## $ GrainYield_Mgha: num [1:1584] 11.4 12.2 12.1 11 10.9 ...
## $ PI_N_Uptake   : num [1:1584] 158 158 158 78 78 ...
```

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

```
paper3_data <- paper3_data %>%
  filter(!ID %in% c(193 , 194 , 721 , 722, 1249 , 1250)) #removes Biggs-18 101 plot bc tractor ran thro
```

```
paper3_data$Platform = factor(paper3_data$Platform, levels=c("GreenSeeker_NDVI" , "sUAS_NDVI" , "sUAS_NDRE
```

```
boxplot(paper3_data$RI ~ paper3_data$Platform)
```



```
paper3_data$RI <- round(paper3_data$RI , digits = 3)
paper3_data$SI <- round(paper3_data$SI , digits = 3)
paper3_data$SI_sq <- round(paper3_data$SI_sq , digits = 3)
```

```
##Data for Table 2
```

```
table_2_data_1 <- paper3_data %>%
  select(site_year , Platform , Index , RI) %>%
  group_by(site_year , Platform) %>%
  summarise(Index_min = min(Index),
            Index_max = max(Index),
            RI_min = min(RI),
            RI_max = max(RI)) %>%
```

```

ungroup()

## `summarise()` has grouped output by 'site_year'. You can override using the `.groups` argument.
table_2_data_1$Index_min <- round(table_2_data_1$Index_min , digits = 2)
table_2_data_1$Index_max <- round(table_2_data_1$Index_max , digits = 2)
table_2_data_1$RI_min <- round(table_2_data_1$RI_min , digits = 2)
table_2_data_1$RI_max <- round(table_2_data_1$RI_max , digits = 2)

table_2_data_2 <- paper3_data %>%
  select(site_year , Platform , PI_N_Uptake) %>%
  group_by(site_year , Platform) %>%
  filter(Platform == "GreenSeeker_NDVI") %>%
  summarise(PI_N_Uptake_min = min(PI_N_Uptake),
            PI_N_Uptake_max = max(PI_N_Uptake)) %>%
  ungroup()

## `summarise()` has grouped output by 'site_year'. You can override using the `.groups` argument.
table_2_data_2$PI_N_Uptake_min <- round(table_2_data_2$PI_N_Uptake_min , digits = 0)
table_2_data_2$PI_N_Uptake_max <- round(table_2_data_2$PI_N_Uptake_max , digits = 0)

#VI MODELS
##GS NDVI
gsndvi_df <- paper3_data %>%
  filter(Platform == "GreenSeeker_NDVI") #filter just gs ndvi values

PI_N_Uptake2 <- gsndvi_df$PI_N_Uptake^2
gs_ndvi_qm <- lm(Index ~ PI_N_Uptake + PI_N_Uptake2 , gsndvi_df) #quadratic model
summary(gs_ndvi_qm) #r2 = 0.80

##
## Call:
## lm(formula = Index ~ PI_N_Uptake + PI_N_Uptake2, data = gsndvi_df)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.182258 -0.040094  0.006751  0.038822  0.207385
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  2.360e-01  1.106e-02   21.34  <2e-16 ***
## PI_N_Uptake   6.169e-03  2.069e-04   29.82  <2e-16 ***
## PI_N_Uptake2 -1.787e-05  8.829e-07  -20.24  <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.05914 on 523 degrees of freedom
## Multiple R-squared:  0.7974, Adjusted R-squared:  0.7967
## F-statistic: 1029 on 2 and 523 DF, p-value: < 2.2e-16

Anova(gs_ndvi_qm , type = 3)

## Anova Table (Type III tests)

```

```
##
## Response: Index
##           Sum Sq   Df F value    Pr(>F)
## (Intercept)  1.5924    1  455.30 < 2.2e-16 ***
## PI_N_Uptake   3.1102    1  889.26 < 2.2e-16 ***
## PI_N_Uptake2  1.4333    1  409.81 < 2.2e-16 ***
## Residuals    1.8292  523
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

gs_ndvi_fit_qm <- fitted(gs_ndvi_qm) #gets fitted values from quad model
gs_ndvi_qm_df <- data.frame(gsndvi_df$PI_N_Uptake , gs_ndvi_fit_qm) #creates dataframe
gs_ndvi_qm_df <- gs_ndvi_qm_df %>%
  mutate(Platform = "GreenSeeker_NDVI")

gs_ndvi_qm_r2 <- round(summary(gs_ndvi_qm)$adj.r.squared , digits = 2)
gs_ndvi_qm_a <- as.numeric(as.character(coef(gs_ndvi_qm)[3]))
gs_ndvi_qm_b <- as.numeric(as.character(coef(gs_ndvi_qm)[2]))
gs_ndvi_qm_c <- as.numeric(as.character(coef(gs_ndvi_qm)[1]))
gs_ndvi_qm_sym_x <- (-gs_ndvi_qm_b) / (2*gs_ndvi_qm_a)
gs_ndvi_qm_sym_y <- gs_ndvi_qm_a*(gs_ndvi_qm_sym_x^2) + gs_ndvi_qm_b*gs_ndvi_qm_sym_x + gs_ndvi_qm_c

gs_ndvi_qm_eqn <- paste("y == -1.8e-05*x^2 + 0.006* x + 0.24")

gs_ndvi_qm_r2

## [1] 0.8
gs_ndvi_qm_sym_y

## [1] 0.7682878
gs_ndvi_qm_sym_x

## [1] 172.5754
##sUAS NDVI
sUASndvi_df <- paper3_data %>%
  filter(Platform == "sUAS_NDVI") #filter just sUAS ndvi values

PI_N_Uptake2 <- sUASndvi_df$PI_N_Uptake^2
sUAS_ndvi_qm <- lm(Index ~ PI_N_Uptake + PI_N_Uptake2 , sUASndvi_df) #quadratic model
summary(sUAS_ndvi_qm) #r2 = 0.78

##
## Call:
## lm(formula = Index ~ PI_N_Uptake + PI_N_Uptake2, data = sUASndvi_df)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.120191 -0.015915  0.000394  0.022040  0.080498
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)   6.696e-01  5.382e-03  124.41  <2e-16 ***
```

```

## PI_N_Uptake    3.268e-03  1.007e-04   32.46   <2e-16 ***
## PI_N_Uptake2 -9.978e-06  4.297e-07  -23.22   <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.02878 on 523 degrees of freedom
## Multiple R-squared:  0.8014, Adjusted R-squared:  0.8006
## F-statistic: 1055 on 2 and 523 DF,  p-value: < 2.2e-16

sUAS_ndvi_fit_qm <- fitted(sUAS_ndvi_qm) #gets fitted values from quad model
sUAS_ndvi_qm_df <- data.frame(sUASndvi_df$PI_N_Uptake , sUAS_ndvi_fit_qm) #creates dataframe

sUAS_ndvi_qm_df <- sUAS_ndvi_qm_df %>%
  mutate(Platform = "sUAS_NDVI")

#getting the equation for the plot
sUAS_ndvi_qm_r2 <- round(summary(sUAS_ndvi_qm)$adj.r.squared , digits = 2)
sUAS_ndvi_qm_a <- as.numeric(as.character(coef(sUAS_ndvi_qm)[3]))
sUAS_ndvi_qm_b <- as.numeric(as.character(coef(sUAS_ndvi_qm)[2]))
sUAS_ndvi_qm_c <- as.numeric(as.character(coef(sUAS_ndvi_qm)[1]))
sUAS_ndvi_qm_sym_x <- (-sUAS_ndvi_qm_b) / (2*sUAS_ndvi_qm_a)
sUAS_ndvi_qm_sym_y <- sUAS_ndvi_qm_a*(sUAS_ndvi_qm_sym_x^2) + sUAS_ndvi_qm_b*sUAS_ndvi_qm_sym_x + sUAS_ndvi_qm_c

sUAS_ndvi_qm_eqn <- paste("y == -1.0e-05*x^2 + 0.0033* x + 0.67")

sUAS_ndvi_qm_r2

## [1] 0.8
sUAS_ndvi_qm_sym_y

## [1] 0.9371271
sUAS_ndvi_qm_sym_x

## [1] 163.7569
##sUAS NDRE

sUASndre_df <- paper3_data %>%
  filter(Platform == "sUAS_NDRE") #filter just sUAS ndre values

PI_N_Uptake2 <- sUASndre_df$PI_N_Uptake^2
sUAS_ndre_qm <- lm(Index ~ PI_N_Uptake + PI_N_Uptake2 , sUASndre_df) #quadratic model
summary(sUAS_ndre_qm) #r2 = 0.82

##
## Call:
## lm(formula = Index ~ PI_N_Uptake + PI_N_Uptake2, data = sUASndre_df)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.141442 -0.034596 -0.001881  0.036173  0.121878
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)   2.738e-01  9.276e-03   29.52   <2e-16 ***

```

```

## PI_N_Uptake    4.442e-03  1.735e-04   25.60   <2e-16 ***
## PI_N_Uptake2 -1.069e-05  7.406e-07  -14.44   <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.04961 on 523 degrees of freedom
## Multiple R-squared:  0.8172, Adjusted R-squared:  0.8165
## F-statistic: 1169 on 2 and 523 DF,  p-value: < 2.2e-16

sUAS_ndre_qm_mse <- mean(residuals(sUAS_ndre_qm)^2)
sUAS_ndre_qm_mse

## [1] 0.002446664

sUAS_ndre_qm_rmse <- sqrt(sUAS_ndre_qm_mse)
sUAS_ndre_qm_rmse

## [1] 0.04946376

sUAS_ndre_fit_qm <- fitted(sUAS_ndre_qm) #gets fitted values from linear model
sUAS_ndre_qm_df <- data.frame(sUASndre_df$PI_N_Uptake , sUAS_ndre_fit_qm) #creates dataframe

sUAS_ndre_qm_df <- sUAS_ndre_qm_df %>%
  mutate(Platform = "sUAS_NDRE")

#getting the equation for the plot
sUAS_ndre_qm_r2 <- round(summary(sUAS_ndre_qm)$adj.r.squared , digits = 2)
sUAS_ndre_qm_a <- as.numeric(as.character(coef(sUAS_ndre_qm)[3]))
sUAS_ndre_qm_b <- as.numeric(as.character(coef(sUAS_ndre_qm)[2]))
sUAS_ndre_qm_c <- as.numeric(as.character(coef(sUAS_ndre_qm)[1]))
sUAS_ndre_qm_sym_x <- (-sUAS_ndre_qm_b) / (2*sUAS_ndre_qm_a)
sUAS_ndre_qm_sym_y <- sUAS_ndre_qm_a*(sUAS_ndre_qm_sym_x^2) + sUAS_ndre_qm_b*sUAS_ndre_qm_sym_x + sUAS_

sUAS_ndre_qm_eqn <- paste("y == -1.1e-05*x^2 + 0.004* x + 0.27")

sUAS_ndre_qm_r2

## [1] 0.82
sUAS_ndre_qm_sym_y

## [1] 0.7351191
sUAS_ndre_qm_sym_x

## [1] 207.7103

#RI MODELS
##GS NDVI-RI

paper3_data_no_2018 <- paper3_data %>%
  filter(year != "2018",
         !ID %in% c(419, 420 , 431 , 432 , 947 , 948 ,959 , 960 , 1475 , 1476 , 1487 , 1488))
#need to get the dataframe in the right form to make saturation figure for the paper. Removes 2018 year

greenseeker_ndvi_data <- paper3_data_no_2018 %>%
  filter(Platform == "GreenSeeker_NDVI")

```

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

gs_ndvi_model <- lme(GrainYield_Mgha ~ SI * TopDress_kgha + I(SI*SI) ,
  control = ctrl ,
  random = ~ I(TopDress_kgha) | site_year ,
  data = greenseeker_ndvi_data)
```

```
summary(gs_ndvi_model)
```

```
## Linear mixed-effects model fit by REML
## Data: greenseeker_ndvi_data
##      AIC      BIC    logLik
## 859.9217 894.0313 -420.9608
##
## Random effects:
## Formula: ~I(TopDress_kgha) | site_year
## Structure: General positive-definite, Log-Cholesky parametrization
##              StdDev   Corr
## (Intercept)  1.0746539 (Intr)
## I(TopDress_kgha) 0.0048649 -0.946
## Residual      0.8137735
##
## Fixed effects: GrainYield_Mgha ~ SI * TopDress_kgha + I(SI * SI)
##              Value Std.Error DF   t-value p-value
## (Intercept)  -0.310935 0.9171844 322 -0.339010  0.7348
## SI           16.478265 2.1839827 322  7.545053  0.0000
## TopDress_kgha  0.083219 0.0134688 322  6.178603  0.0000
## I(SI * SI)    -5.517499 1.4660309 322 -3.763563  0.0002
## SI:TopDress_kgha -0.079362 0.0150575 322 -5.270580  0.0000
## Correlation:
##              (Intr) SI      TpDrs_ I(SI*S
## SI           -0.842
## TopDress_kgha -0.311  0.080
## I(SI * SI)    0.781 -0.982  0.055
## SI:TopDress_kgha 0.242 -0.083 -0.973 -0.056
##
## Standardized Within-Group Residuals:
##      Min      Q1      Med      Q3      Max
## -3.63978666 -0.56804670  0.03176909  0.63982598  3.32828753
##
## Number of Observations: 332
## Number of Groups: 6
```

```
summary(gs_ndvi_model)$tTable
```

```
##              Value Std.Error DF   t-value    p-value
## (Intercept)  -0.31093457 0.91718440 322 -0.3390099 7.348233e-01
## SI           16.47826480 2.18398272 322  7.5450527 4.668881e-13
## TopDress_kgha  0.08321852 0.01346883 322  6.1786031 1.951965e-09
## I(SI * SI)    -5.51749911 1.46603094 322 -3.7635625 1.990236e-04
## SI:TopDress_kgha -0.07936173 0.01505749 322 -5.2705803 2.496678e-07
```

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

```
## Analysis of Deviance Table (Type III tests)
```



```
##
## Response: GrainYield_Mgha
##               Chisq Df Pr(>Chisq)
## (Intercept)    0.1149  1  0.7346023
## SI             56.9278  1  4.521e-14 ***
## TopDress_kgha  38.1751  1  6.467e-10 ***
## I(SI * SI)     14.1644  1  0.0001675 ***
## SI:TopDress_kgha 27.7790  1  1.360e-07 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

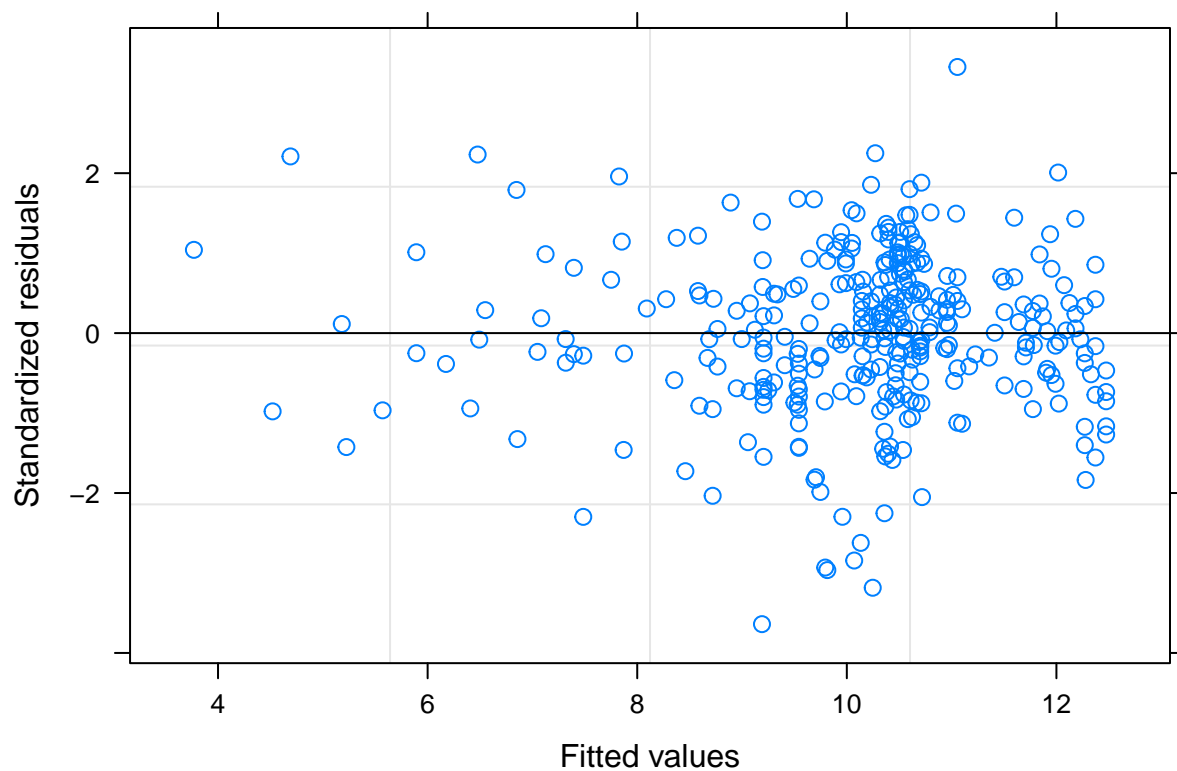
```
r.squaredGLMM(gs_ndvi_model)
```

```
## Warning: 'r.squaredGLMM' now calculates a revised statistic. See the help page.
```

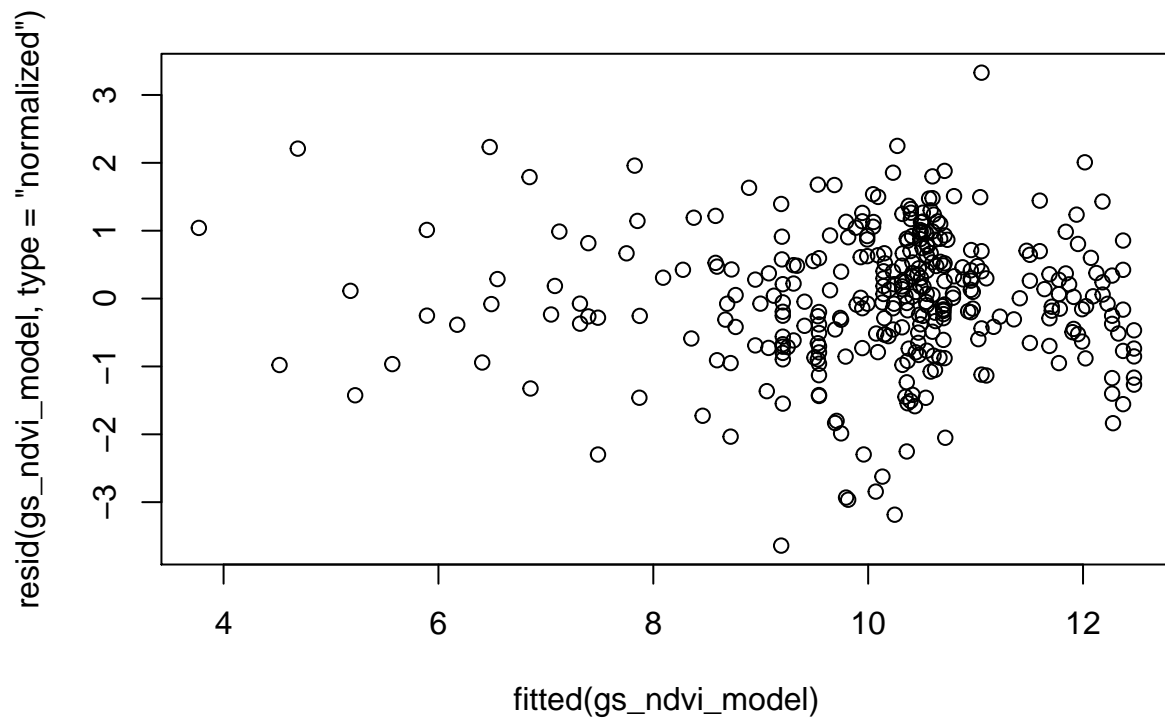
```
##           R2m      R2c
## [1,] 0.4483278 0.776302
```

model diagnostics

```
plot(gs_ndvi_model)
```



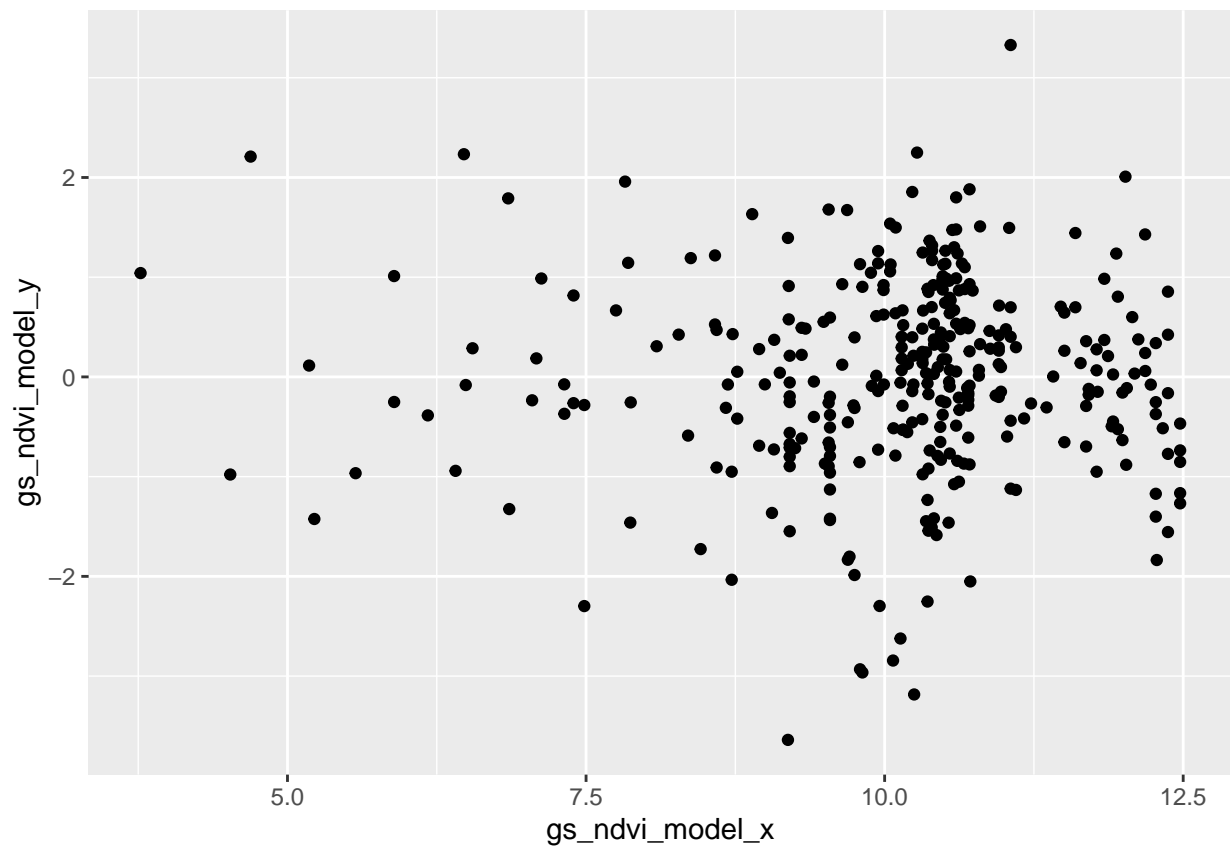
```
plot(resid(gs_ndvi_model, type = "normalized") ~fitted(gs_ndvi_model))
```



```
gs_ndvi_model_y <- resid(gs_ndvi_model, type = "normalized")
gs_ndvi_model_x <- fitted(gs_ndvi_model)

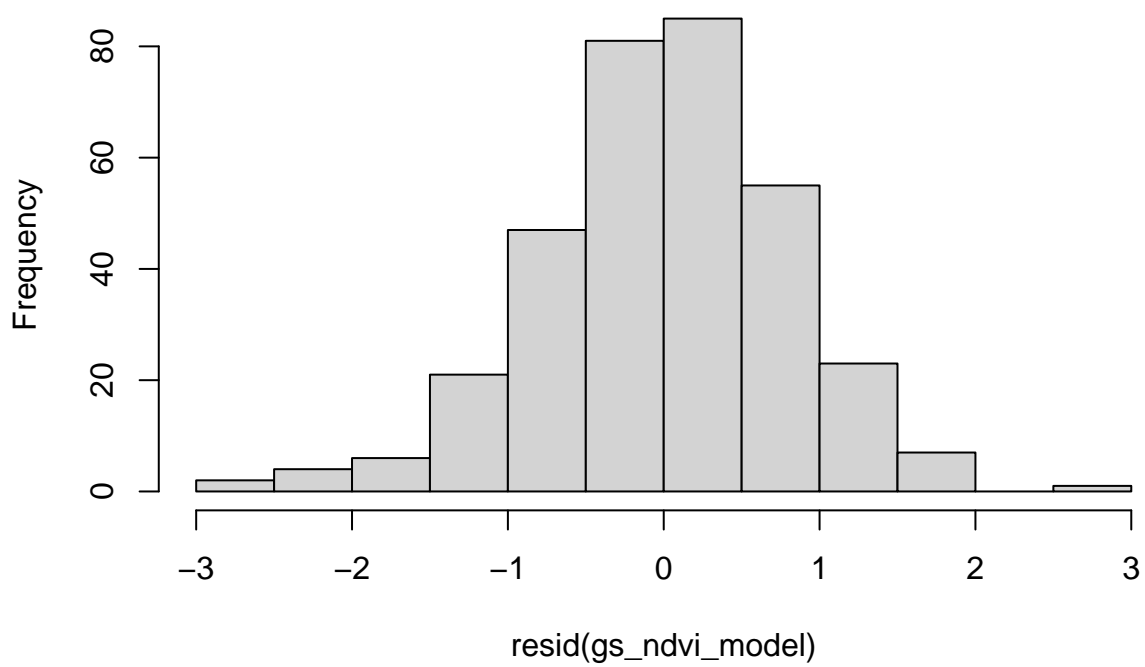
gs_ndvi_modelresid_data <- data.frame(gs_ndvi_model_x , gs_ndvi_model_y)

ggplot( data = gs_ndvi_modelresid_data , aes( x = gs_ndvi_model_x , y = gs_ndvi_model_y)) +
  geom_point(mapping = aes(gs_ndvi_model_x , gs_ndvi_model_y) , data = gs_ndvi_modelresid_data)
```

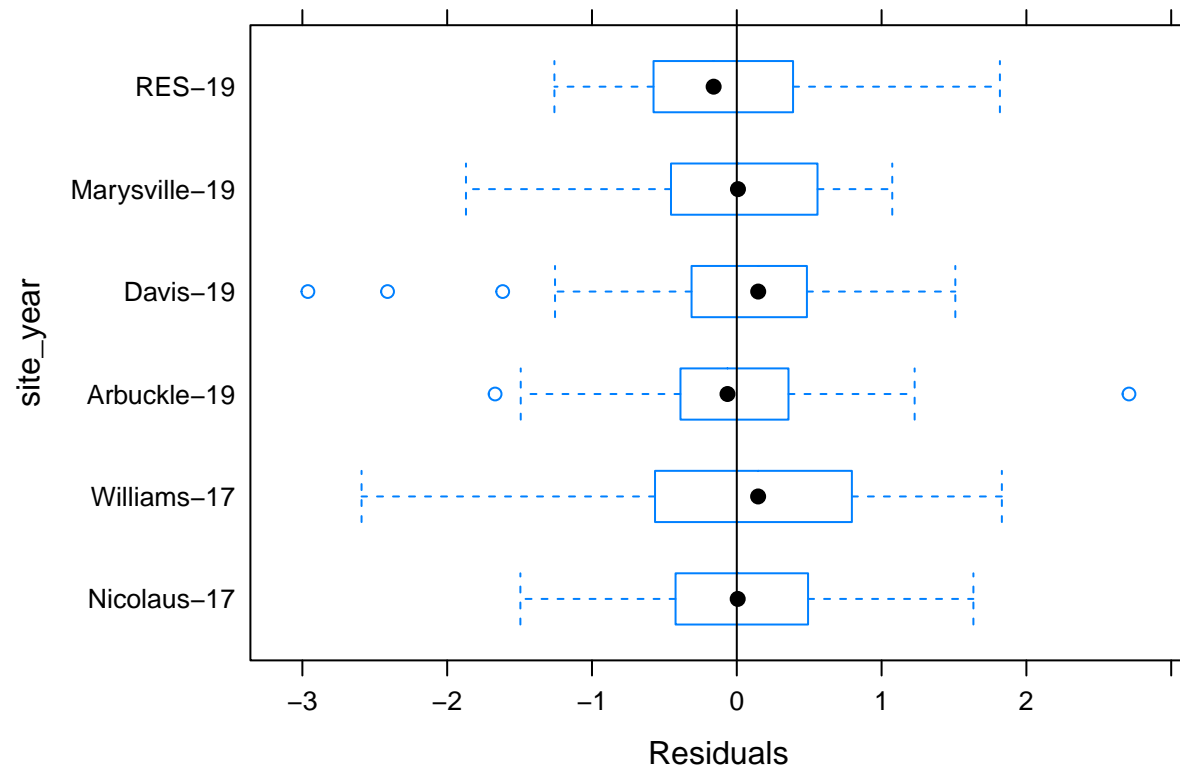


```
hist(resid(gs_ndvi_model))
```

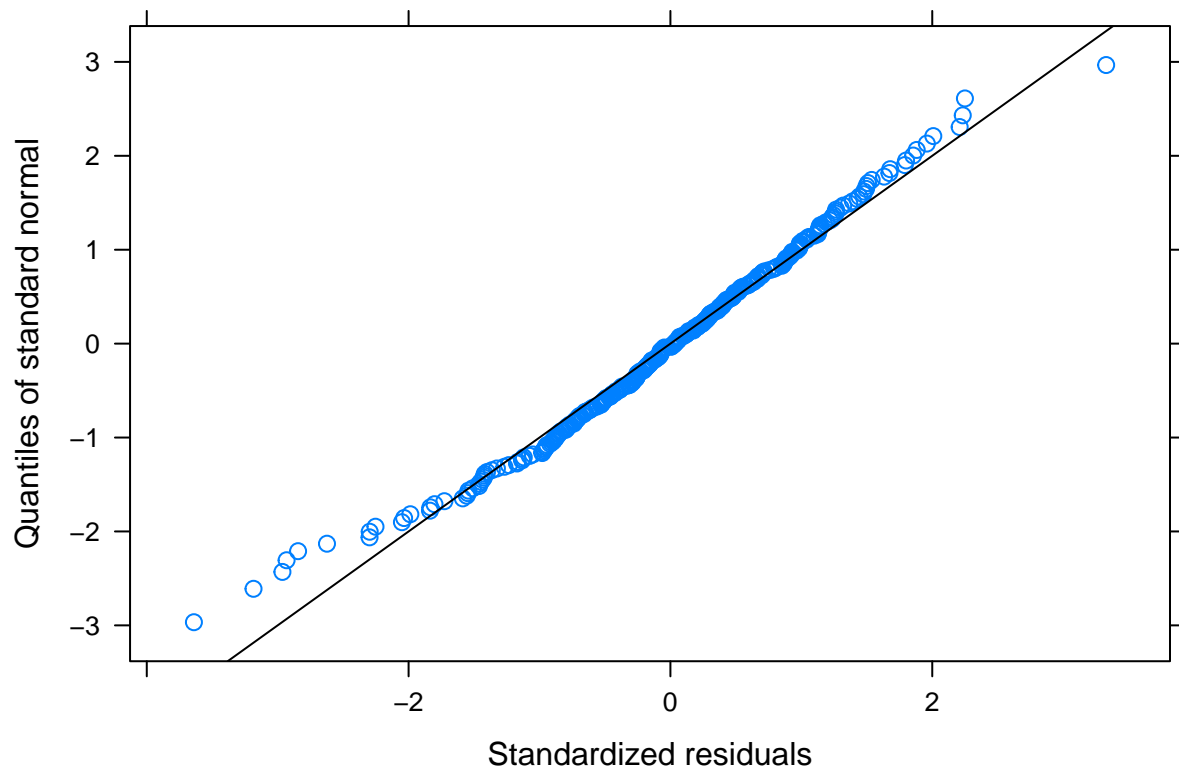
Histogram of resid(gs_ndvi_model)



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

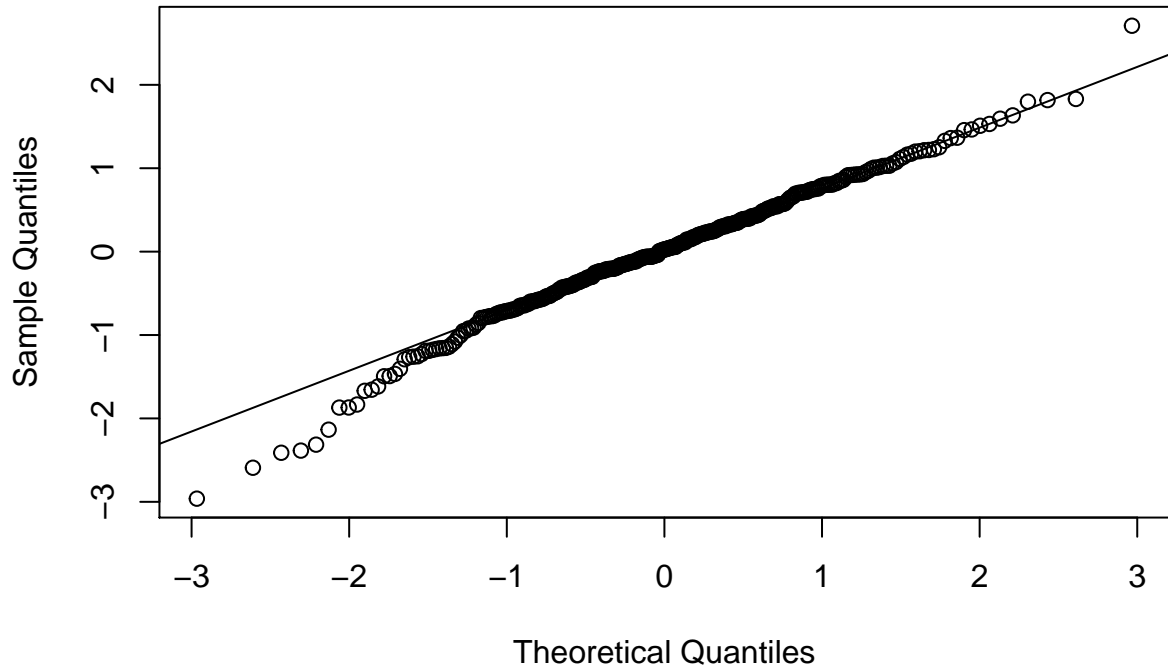


```
qqnorm(gs_ndvi_model, abline = c(0,1) )
```

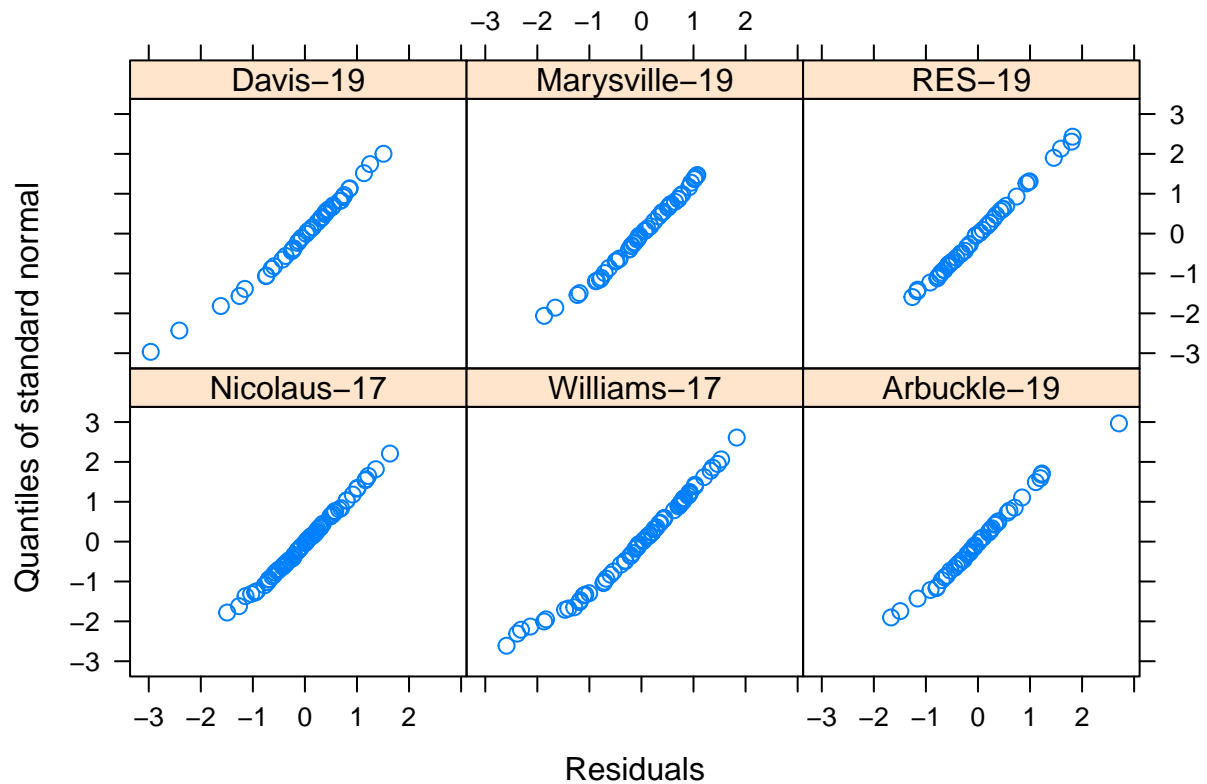


```
qqnorm(resid(gs_ndvi_model))
qqline(resid(gs_ndvi_model))
```

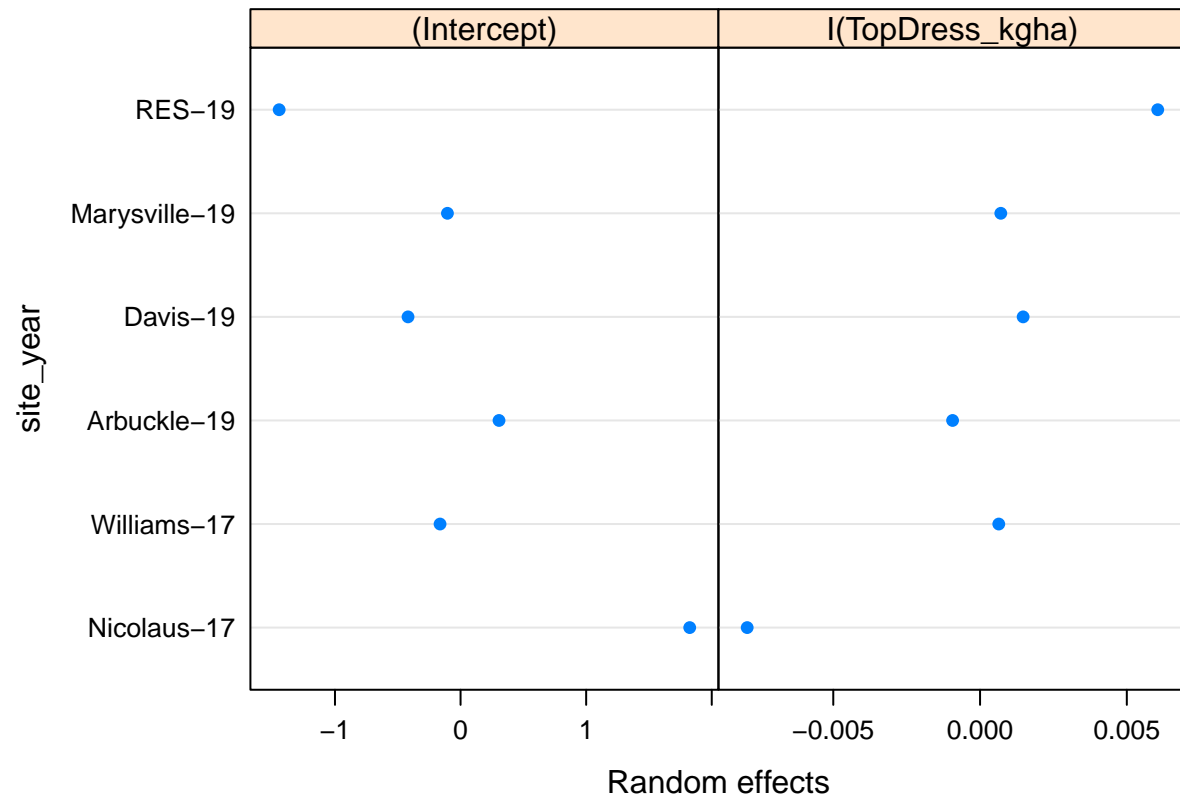
Normal Q-Q Plot



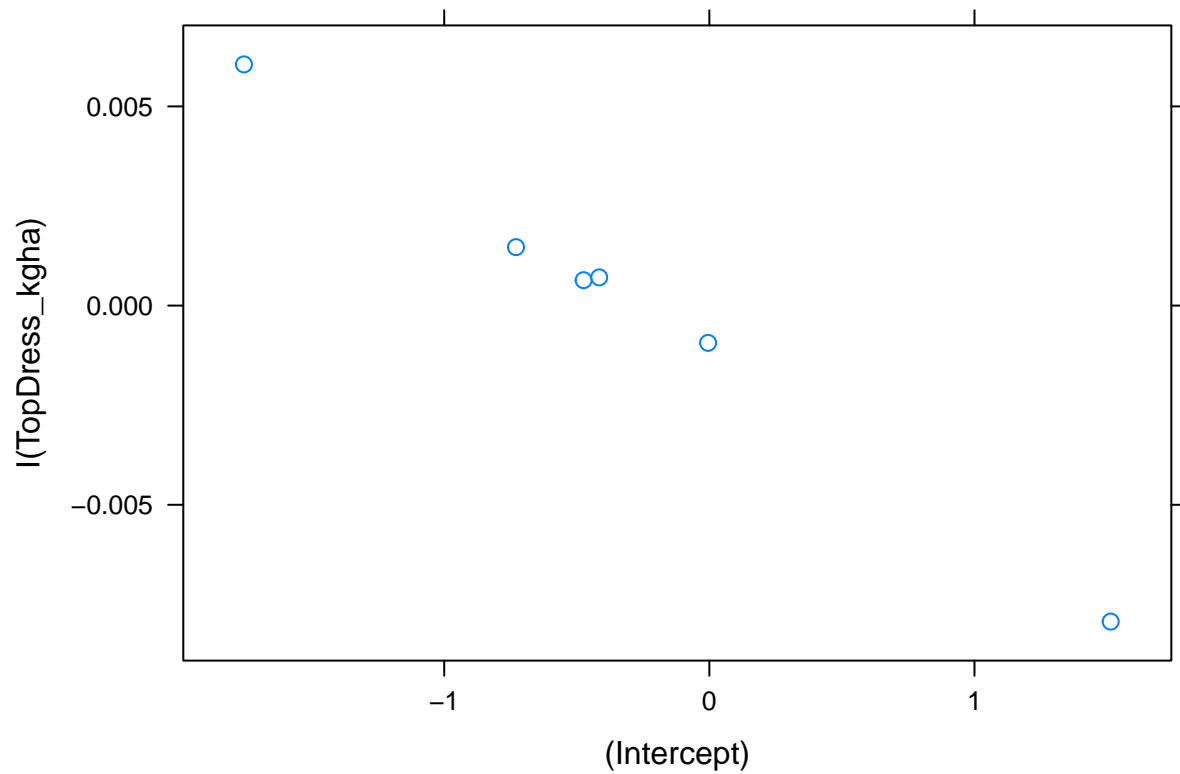
```
qqnorm(gs_ndvi_model , ~resid(.) | site_year)
```



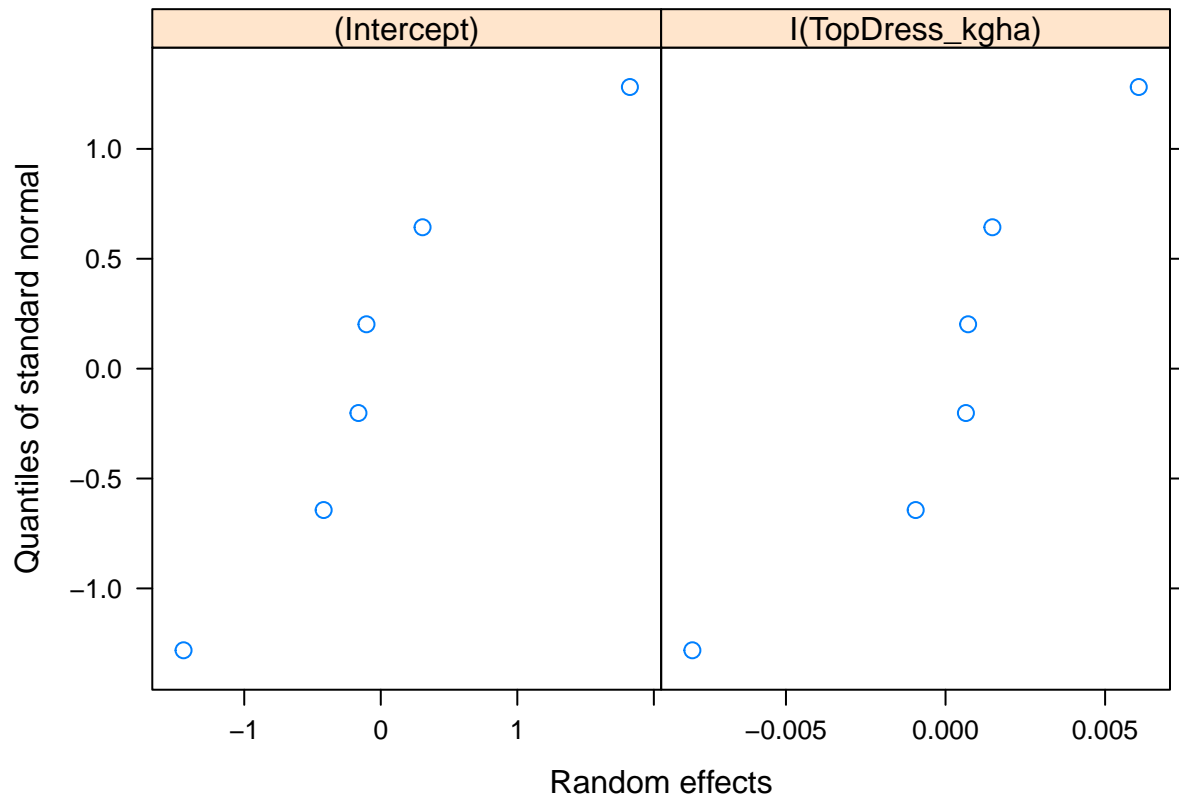
```
plot(ranef(gs_ndvi_model))
```



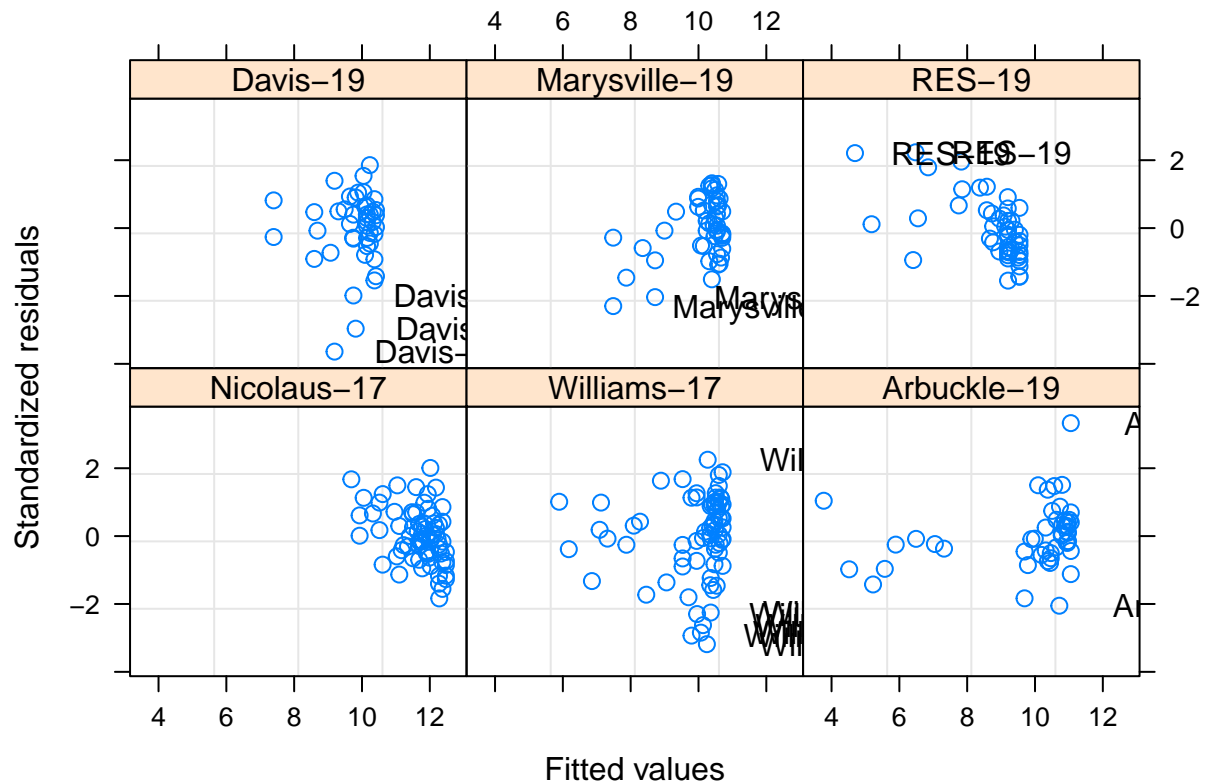
```
pairs(gs_ndvi_model , id = 0.1)
```



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



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



```
###outlier removal
```

```
greenseeker_ndvi_data_OR <- greenseeker_ndvi_data[-(c(which(abs(residuals(gs_ndvi_model,type="normalized
```

```
###rerun model sans outliers
```

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

```
gs_ndvi_model_OR <- lme(GrainYield_Mgha ~ SI * TopDress_kgha + I(SI*SI) ,
  control = ctrl ,
  random = ~ I(TopDress_kgha) | site_year ,
  data = greenseeker_ndvi_data_OR)
```

```
summary(gs_ndvi_model_OR)
```

```
## Linear mixed-effects model fit by REML
```

```
## Data: greenseeker_ndvi_data_OR
```

```
## AIC BIC logLik
```

```
## 818.3621 852.3888 -400.1811
```

```
##
```

```
## Random effects:
```

```
## Formula: ~I(TopDress_kgha) | site_year
```

```
## Structure: General positive-definite, Log-Cholesky parametrization
```

```
## StdDev Corr
```

```
## (Intercept) 1.061801398 (Intr)
```

```
## I(TopDress_kgha) 0.004744882 -0.942
```

```
## Residual 0.771366042
```

```
##
```

```
## Fixed effects: GrainYield_Mgha ~ SI * TopDress_kgha + I(SI * SI)
```

```
## Value Std.Error DF t-value p-value
```



```
## (Intercept)      -0.476590 0.8784971 319 -0.542506 0.5878
## SI               17.147163 2.0736291 319  8.269156 0.0000
## TopDress_kgha    0.082491 0.0127849 319  6.452271 0.0000
## I(SI * SI)       -6.015697 1.3924269 319 -4.320296 0.0000
## SI:TopDress_kgha -0.079565 0.0142834 319 -5.570456 0.0000
## Correlation:
##               (Intr) SI      TpDrs_ I(SI*S
## SI            -0.834
## TopDress_kgha -0.311 0.078
## I(SI * SI)     0.774 -0.982 0.056
## SI:TopDress_kgha 0.239 -0.082 -0.972 -0.056
##
## Standardized Within-Group Residuals:
##      Min      Q1      Med      Q3      Max
## -3.21293731 -0.58644957 0.03185367 0.66072053 2.35677933
##
## Number of Observations: 329
## Number of Groups: 6
```

```
summary(gs_ndvi_model_OR)$tTable
```

```
##              Value Std.Error DF   t-value    p-value
## (Intercept)  -0.47658960 0.87849707 319 -0.5425056 5.878490e-01
## SI           17.14716306 2.07362910 319  8.2691563 3.690901e-15
## TopDress_kgha  0.08249138 0.01278486 319  6.4522706 4.088965e-10
## I(SI * SI)     -6.01569684 1.39242688 319 -4.3202964 2.083895e-05
## SI:TopDress_kgha -0.07956498 0.01428339 319 -5.5704563 5.404405e-08
```

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

```
## Analysis of Deviance Table (Type III tests)
##
## Response: GrainYield_Mgha
##              Chisq Df Pr(>Chisq)
## (Intercept)    0.2943 1    0.5875
## SI             68.3789 1 < 2.2e-16 ***
## TopDress_kgha  41.6318 1 1.102e-10 ***
## I(SI * SI)     18.6650 1 1.558e-05 ***
## SI:TopDress_kgha 31.0300 1 2.541e-08 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
r.squaredGLMM(gs_ndvi_model_OR)
```

```
##              R2m      R2c
## [1,] 0.4590344 0.7917617
```

```
gs_ndvi_r_sq <- r.squaredGLMM(gs_ndvi_model_OR)
```

```
gs_ndvi_r_sq_fixed <- round(gs_ndvi_r_sq[1] , digits = 2)
gs_ndvi_r_sq_fixed
```

```
## [1] 0.46
```

```
gs_ndvi_r_sq_total <- round(gs_ndvi_r_sq[2] , digits = 2)
gs_ndvi_r_sq_total
```

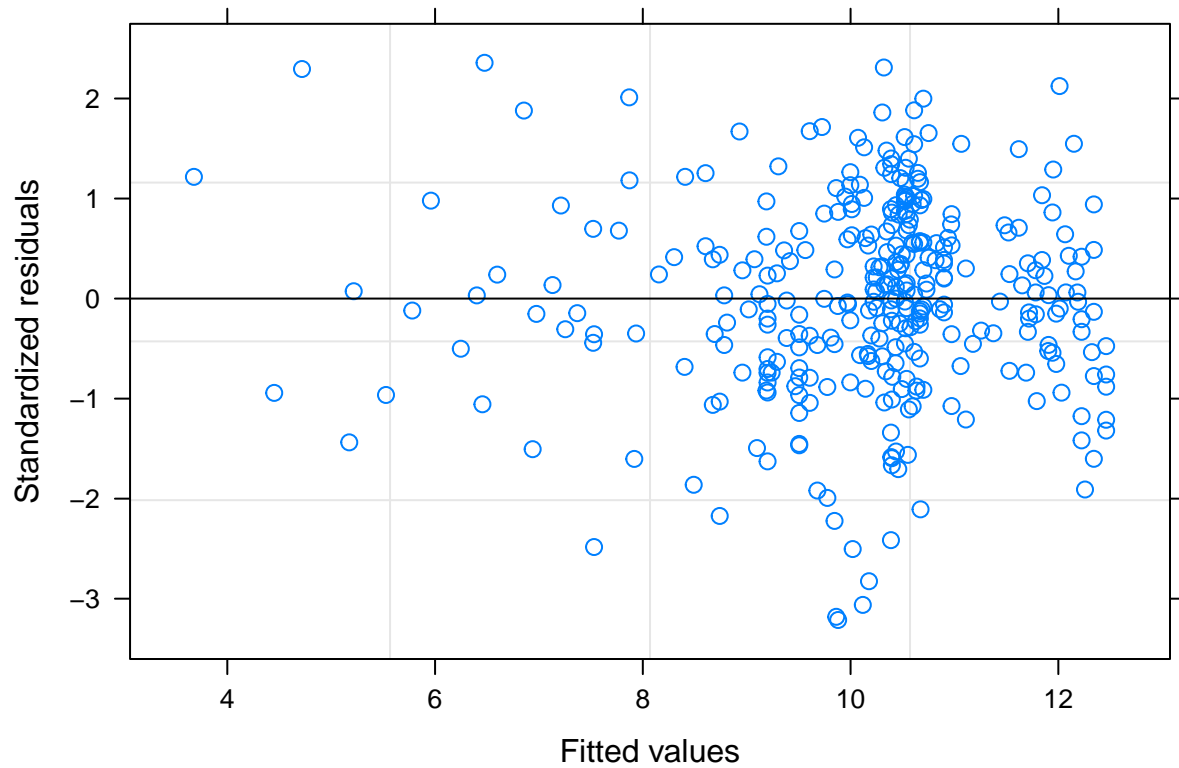
```
## [1] 0.79
```

```
gs_ndvi_r_sq_random <- gs_ndvi_r_sq_total - gs_ndvi_r_sq_fixed  
gs_ndvi_r_sq_random
```

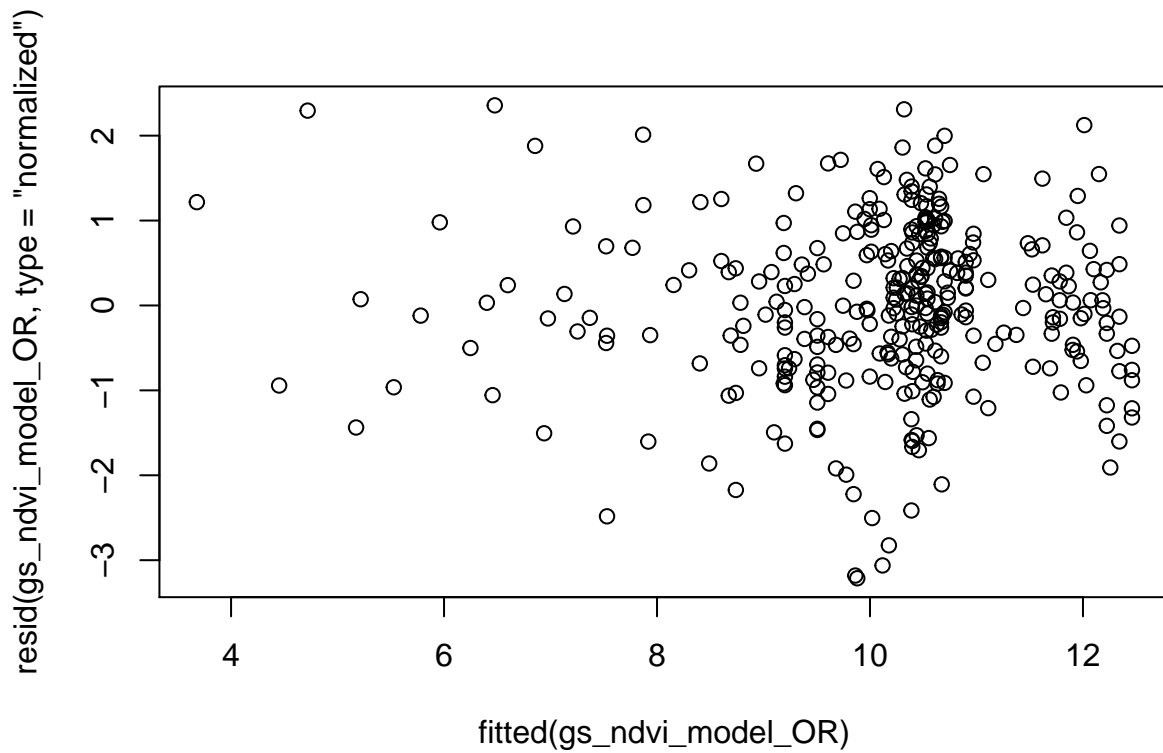
```
## [1] 0.33
```

model diagnostics

```
plot (gs_ndvi_model_OR)
```



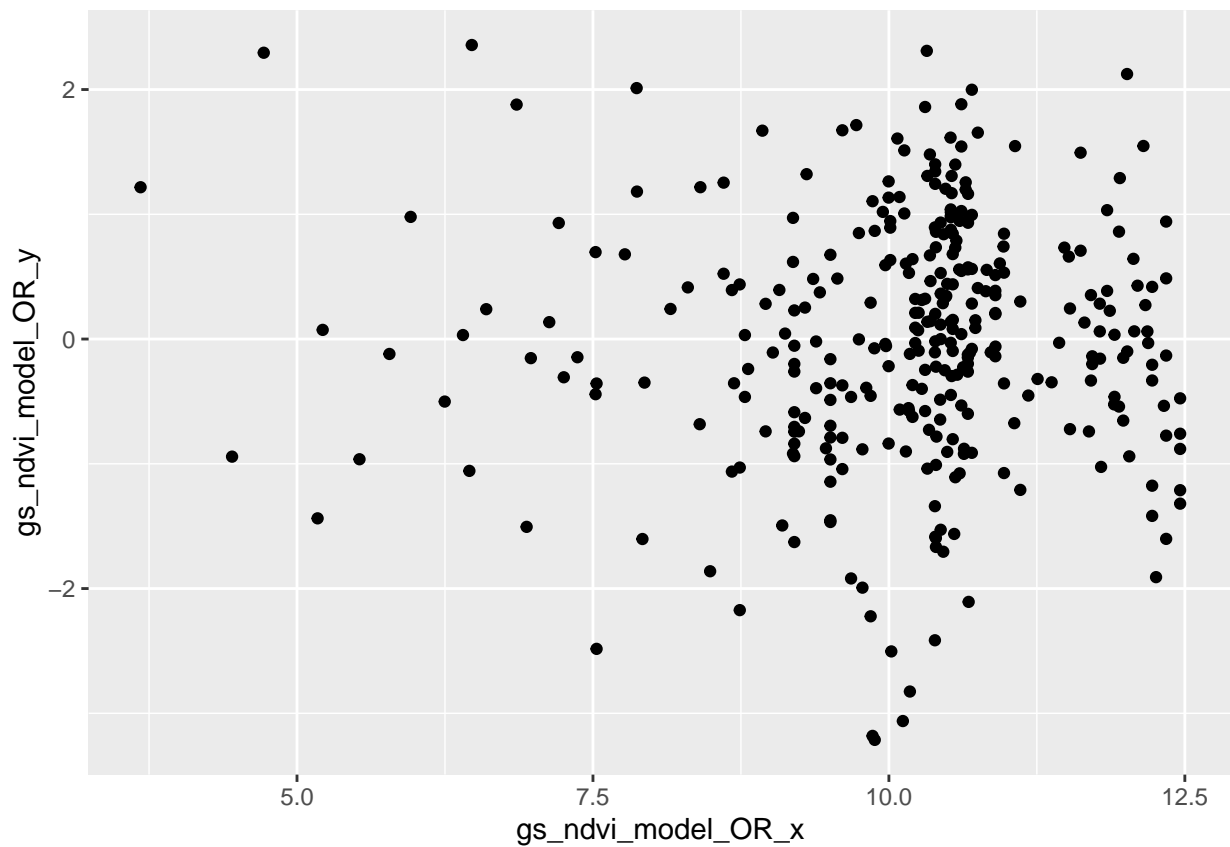
```
plot(resid(gs_ndvi_model_OR, type = "normalized") ~fitted(gs_ndvi_model_OR))
```



```
gs_ndvi_model_OR_y <- resid(gs_ndvi_model_OR, type = "normalized")
gs_ndvi_model_OR_x <- fitted(gs_ndvi_model_OR)

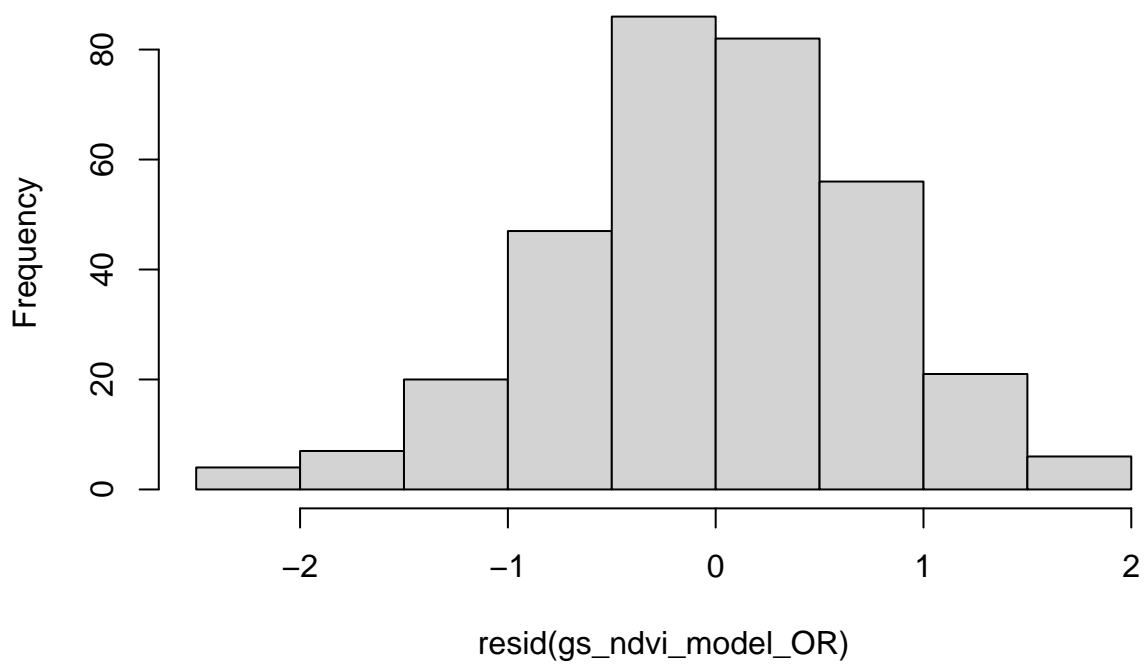
gs_ndvi_model_ORresid_data <- data.frame(gs_ndvi_model_OR_x , gs_ndvi_model_OR_y)

ggplot( data = gs_ndvi_model_ORresid_data , aes( x = gs_ndvi_model_OR_x , y = gs_ndvi_model_OR_y)) +
  geom_point(mapping = aes(gs_ndvi_model_OR_x , gs_ndvi_model_OR_y) , data = gs_ndvi_model_ORresid_data)
```

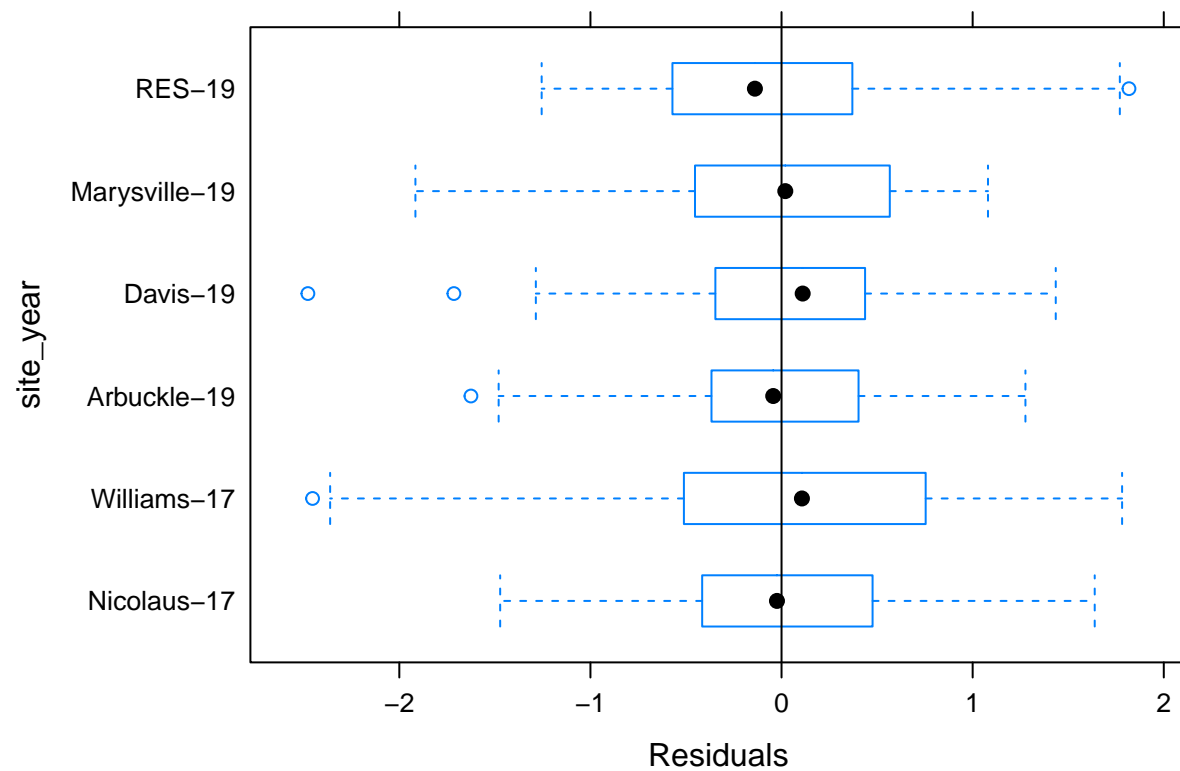


```
hist(resid(gs_ndvi_model_OR))
```

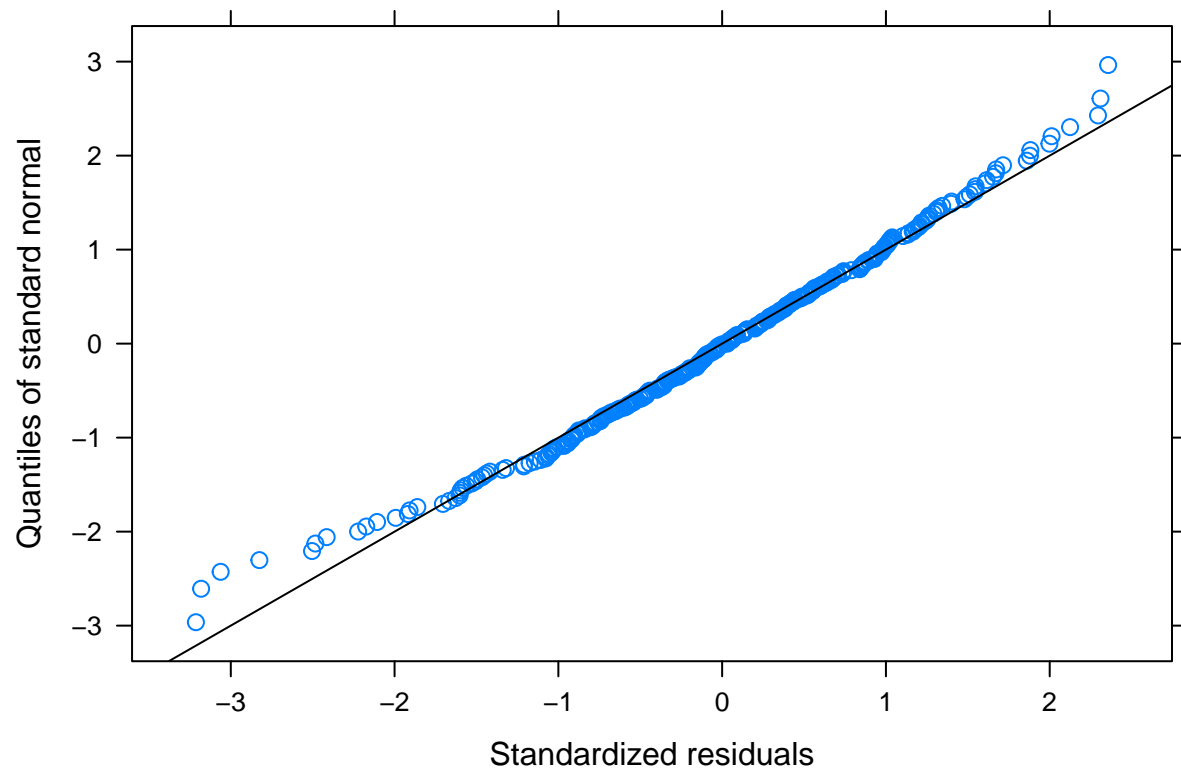
Histogram of resid(gs_ndvi_model_OR)



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

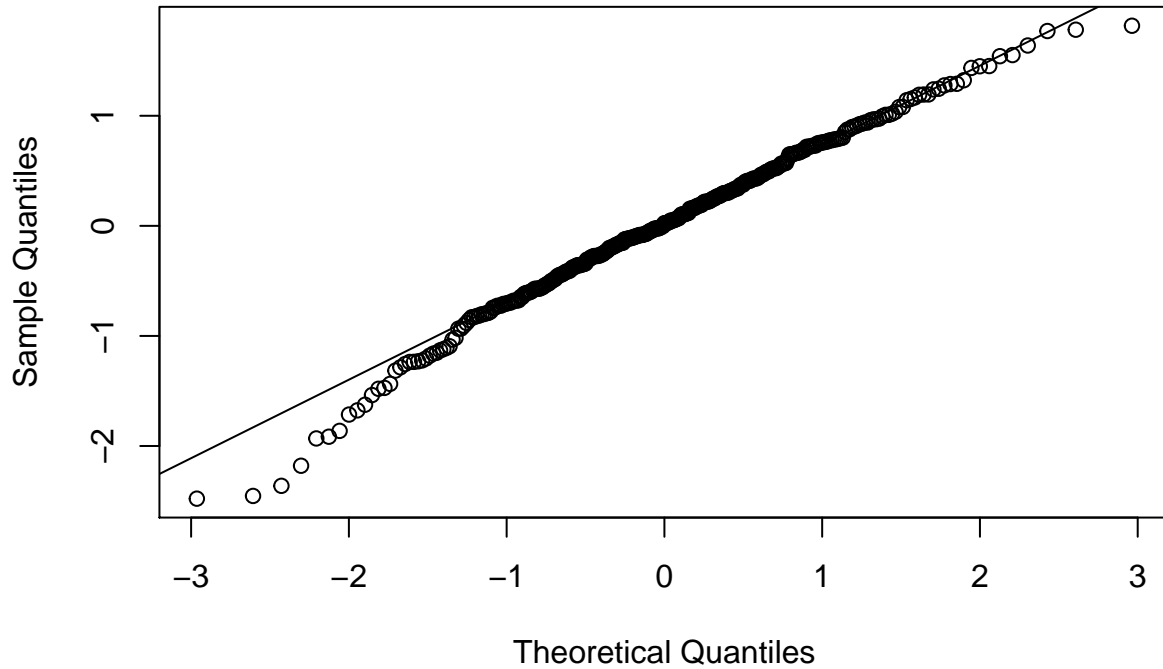


```
qqnorm(gs_ndvi_model_OR, abline = c(0,1) )
```

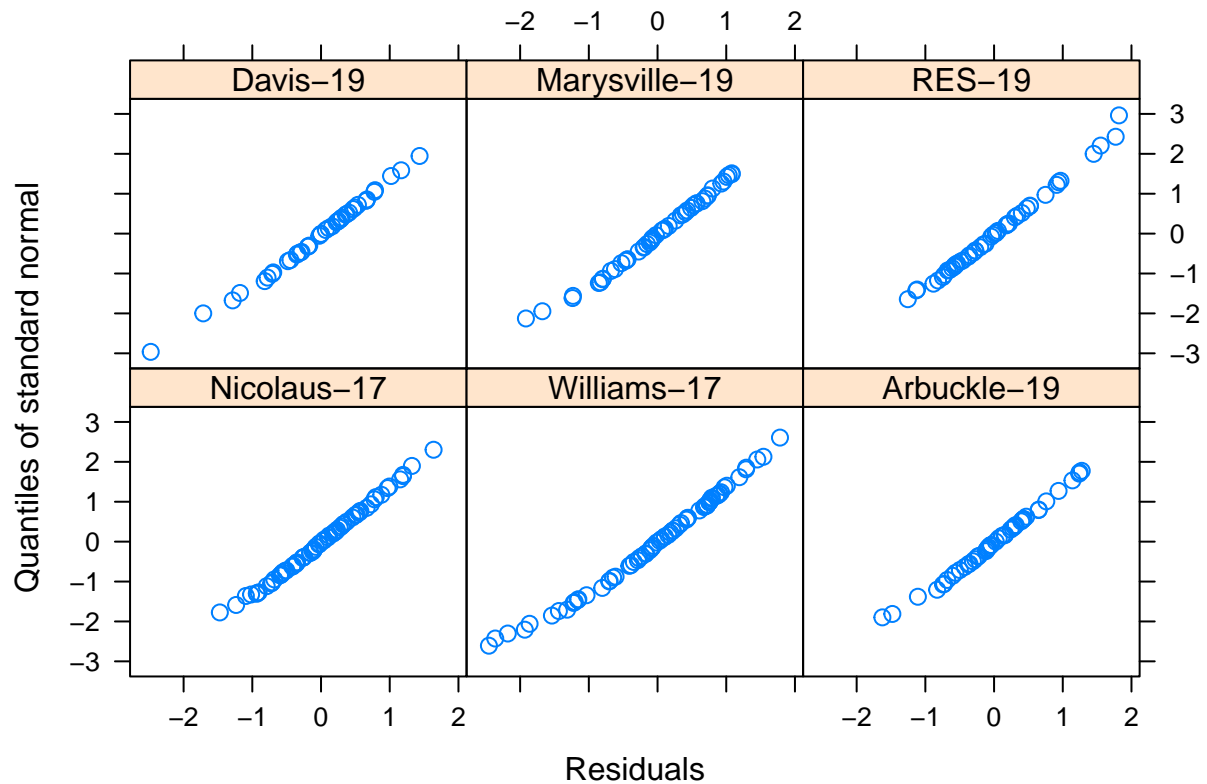


```
qqnorm(resid(gs_ndvi_model_OR))
qqline(resid(gs_ndvi_model_OR))
```

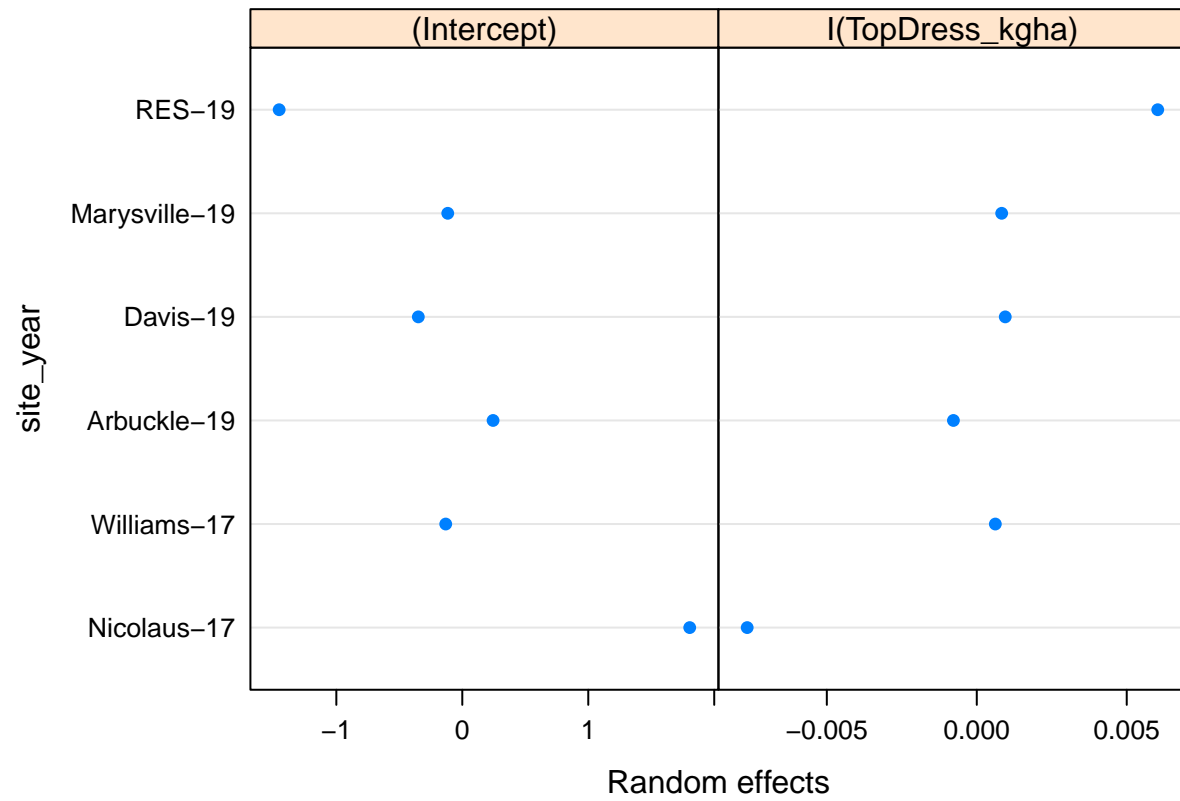
Normal Q-Q Plot



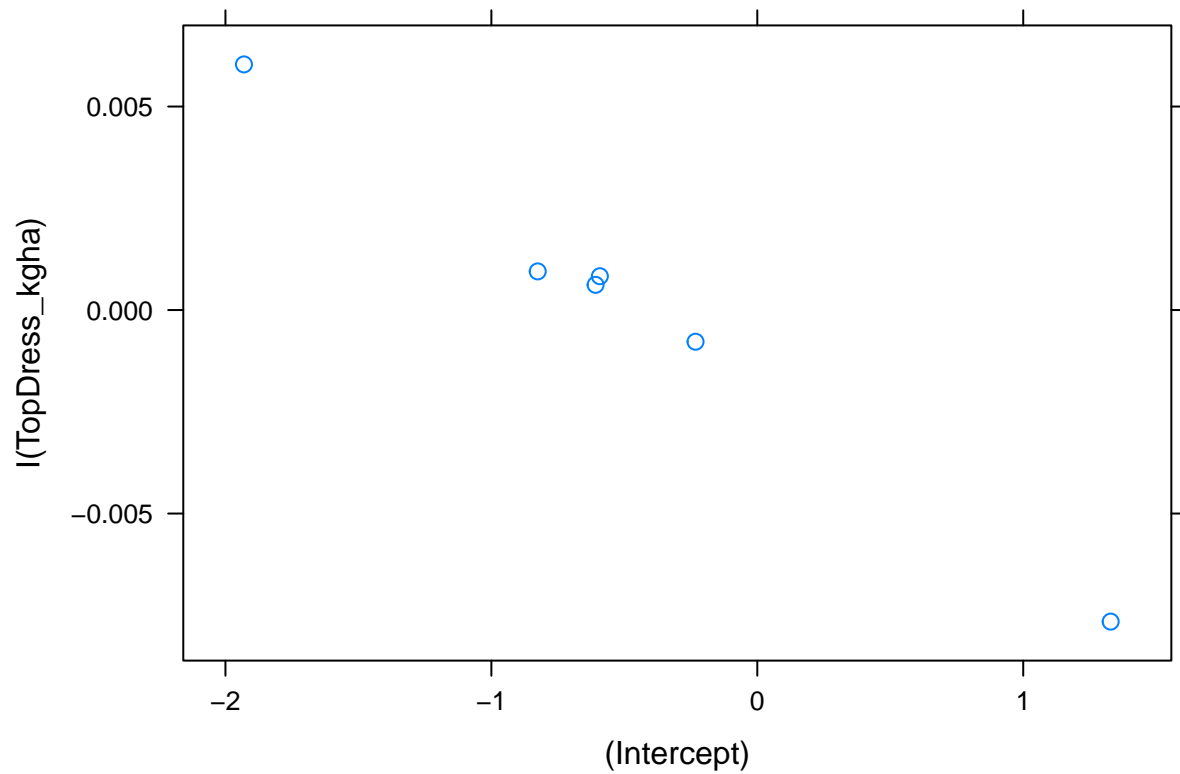
```
qqnorm(gs_ndvi_model_OR , ~resid(.) | site_year)
```



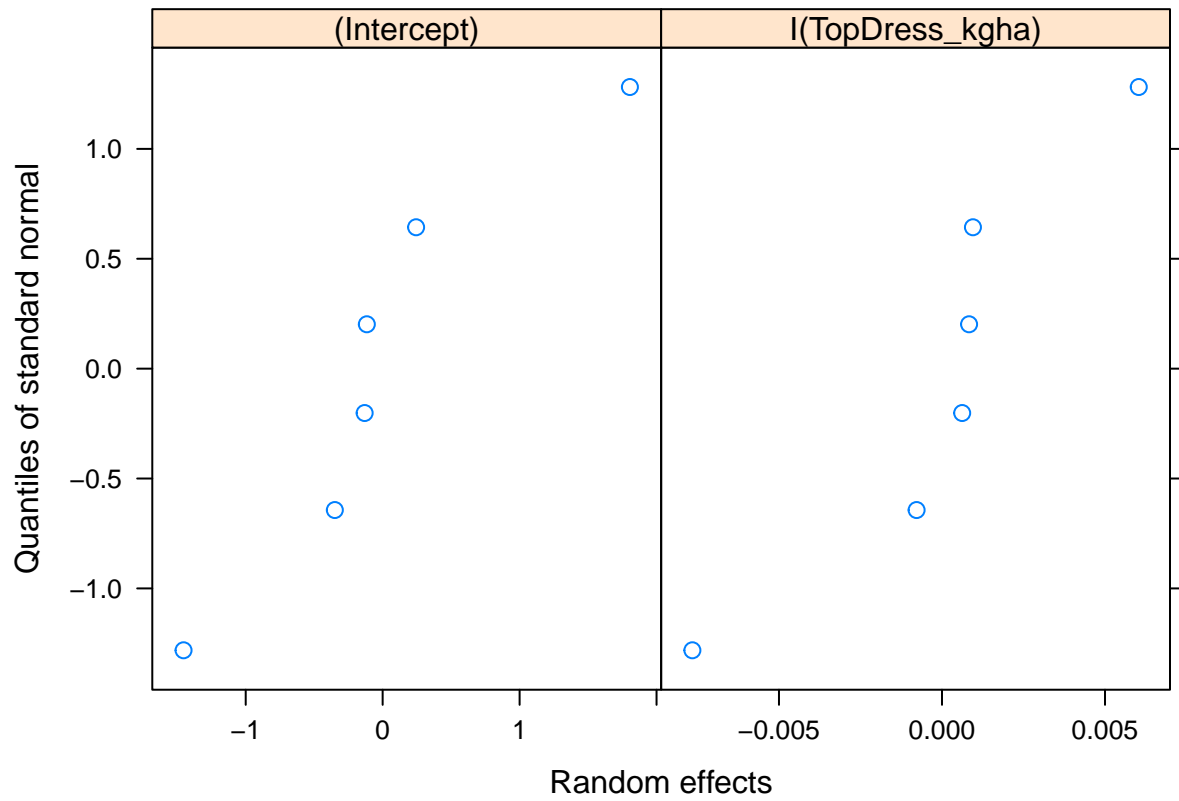
```
plot(ranef(gs_ndvi_model_OR))
```



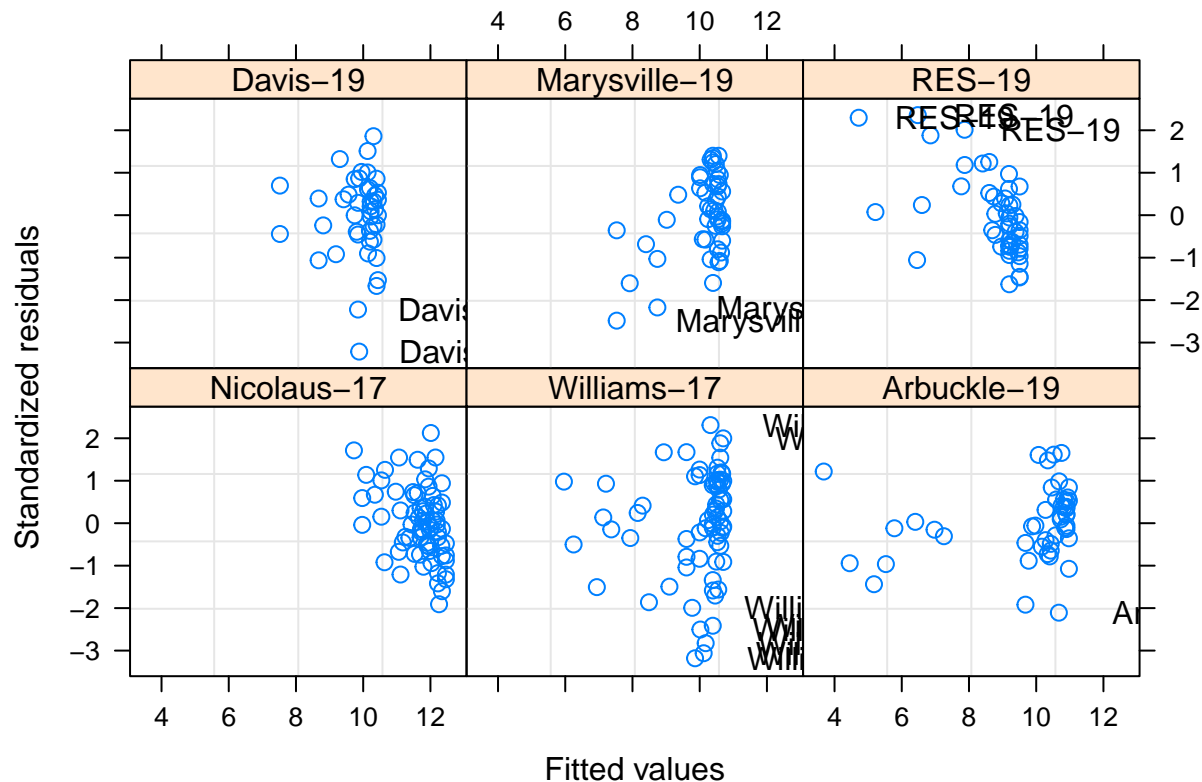
```
pairs(gs_ndvi_model_OR , id = 0.1)
```



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



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

```
### emmeans
```

```
mylist <- list(SI=seq(0.25 , 1 , by = .0005), TopDress_kgha =c( 34 , 0))

gs_ndvi_emmeans <- emmeans(gs_ndvi_model_OR , ~TopDress_kgha * SI , at = mylist )

gs_ndvi_emmeans_contrast <- as.data.frame(summary(contrast(gs_ndvi_emmeans , "pairwise" , side = ">" , l

gs_ndvi_emmeans_contrast <- gs_ndvi_emmeans_contrast %>%
  mutate(GS_NDVI_Response_Index = 1 / SI )

gs_ndvi_emmeans_contrast <- gs_ndvi_emmeans_contrast %>%
  mutate(prob_postive_resp = (1 - p.value) * 100)

gs_ndvi_emmeans_contrast <- gs_ndvi_emmeans_contrast %>%
  mutate(t_score = abs((estimate - 0.26)) / SE)

gs_ndvi_emmeans_contrast <- gs_ndvi_emmeans_contrast %>%
  mutate(prob = if_else(estimate < 0.26 , pt(q = t_score , df = df , lower.tail = F) , pt(q = t_score ,

gs_ndvi_emmeans_contrast$prob_percent <- (round(gs_ndvi_emmeans_contrast$prob , digits = 3) * 100)

gs_ndvi_emmeans_contrast$GS_NDVI_Response_Index_r <- round(gs_ndvi_emmeans_contrast$GS_NDVI_Response_In

gs_ndvi_run1 <- 125 - 100
gs_ndvi_rise1 <- 0.6404977 - 0.09953884
gs_ndvi_slope1 <- round((gs_ndvi_rise1 / gs_ndvi_run1) * 5 , digits = 2)
gs_ndvi_slope1
```

```

## [1] 0.11
gs_ndvi_run2 <- 250 - 125
gs_ndvi_rise2 <- 1.722415 - 0.6404977
gs_ndvi_slope2 <- round((gs_ndvi_rise2 / gs_ndvi_run2) * 5 , digits = 3)
gs_ndvi_slope2

## [1] 0.043
gs_ndvi_run3 <- 400 - 250
gs_ndvi_rise3 <- 2.128135 - 1.722415
gs_ndvi_slope3 <- round((gs_ndvi_rise3 / gs_ndvi_run3) * 5 , digits = 3)
gs_ndvi_slope3

## [1] 0.014
gs_ndvi_mean_se <- gs_ndvi_emmeans_contrast %>%
  select(SE) %>%
  summarise(mean_se = mean(SE))

gs_ndvi_mean_se$mean_se <- round(gs_ndvi_mean_se$mean_se , digits = 2)

gs_ndvi_mean_se$mean_se

## [1] 0.18
###confidence interval
#at every 0.10 interval, what is the 90% confidence interval of the yield response, and what RI value d

str(gs_ndvi_emmeans_contrast , give.attr = F)

## 'data.frame': 1501 obs. of 13 variables:
## $ contrast : Factor w/ 1 level "34 - 0": 1 1 1 1 1 1 1 1 1 1 ...
## $ SI : num 0.25 0.251 0.251 0.252 0.252 ...
## $ estimate : num 2.13 2.13 2.13 2.12 2.12 ...
## $ SE : num 0.318 0.318 0.317 0.317 0.317 ...
## $ df : num 319 319 319 319 319 319 319 319 319 319 ...
## $ t.ratio : num 6.69 6.69 6.7 6.7 6.7 ...
## $ p.value : num 4.88e-11 4.87e-11 4.85e-11 4.83e-11 4.82e-11 ...
## $ GS_NDVI_Response_Index : num 4 3.99 3.98 3.98 3.97 ...
## $ prob_postive_resp : num 100 100 100 100 100 ...
## $ t_score : num 5.88 5.88 5.88 5.88 5.88 ...
## $ prob : num 1 1 1 1 1 ...
## $ prob_percent : num 100 100 100 100 100 100 100 100 100 100 ...
## $ GS_NDVI_Response_Index_r: num 4 3.99 3.98 3.98 3.97 ...

gs_ndvi_emmeans_contrast <- gs_ndvi_emmeans_contrast %>%
  mutate(response_mgha_r = round(estimate , digits = 2),
         se_mgha_r = round(SE , digits = 3))

z_score <- 1.645

#estimate = 0.10 , SE = 0.120

print(round(0.10 - z_score*0.120 , digits = 2))

## [1] -0.1

```

```

print(round(0.10 + z_score*0.120 , digits = 2))

## [1] 0.3
#estimate = 0.20 , SE = 0.112

print(round(0.20 - z_score*0.112 , digits = 2))

## [1] 0.02
print(round(0.20 + z_score*0.112 , digits = 2))

## [1] 0.38
#estimate = 0.30 , SE = 0.106

print(round(0.30 - z_score*0.106 , digits = 2))

## [1] 0.13
print(round(0.30 + z_score*0.106 , digits = 2))

## [1] 0.47
#estimate = 0.40 , SE = 0.102

print(round(0.40 - z_score*0.102 , digits = 2))

## [1] 0.23
print(round(0.40 + z_score*0.102 , digits = 2))

## [1] 0.57
#estimate = 0.50 , SE = 0.102

print(round(0.50 - z_score*0.102 , digits = 2))

## [1] 0.33
print(round(0.50 + z_score*0.102 , digits = 2))

## [1] 0.67
##sUAS NDVI-RI

sUAS_ndvi_data <- paper3_data_no_2018 %>%
  filter(Platform == "sUAS_NDVI")

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

sUAS_ndvi_model <- lme(GrainYield_Mgha ~ SI * TopDress_kgha + I(SI*SI) ,
  control = ctrl ,
  random = ~ I(TopDress_kgha) | site_year ,
  data = sUAS_ndvi_data)

summary(sUAS_ndvi_model)

## Linear mixed-effects model fit by REML
##   Data: sUAS_ndvi_data
##       AIC       BIC    logLik

```

```
##      821.447 855.5566 -401.7235
##
## Random effects:
## Formula: ~I(TopDress_kgha) | site_year
## Structure: General positive-definite, Log-Cholesky parametrization
##              StdDev      Corr
## (Intercept)    1.071047781 (Intr)
## I(TopDress_kgha) 0.004528693 -0.935
## Residual        0.776505570
##
## Fixed effects: GrainYield_Mgha ~ SI * TopDress_kgha + I(SI * SI)
##              Value Std.Error DF   t-value p-value
## (Intercept)   -24.39218  7.976495 322 -3.058007  0.0024
## SI             47.63446 17.706195 322  2.690271  0.0075
## TopDress_kgha   0.20859  0.042869 322  4.865671  0.0000
## I(SI * SI)     -12.53880  9.815347 322 -1.277468  0.2024
## SI:TopDress_kgha -0.20263  0.044502 322 -4.553251  0.0000
## Correlation:
##              (Intr) SI      TpDrs_ I(SI*S
## SI            -0.996
## TopDress_kgha -0.025 -0.031
## I(SI * SI)     0.989 -0.998  0.078
## SI:TopDress_kgha 0.023  0.031 -0.998 -0.078
##
## Standardized Within-Group Residuals:
##              Min          Q1          Med          Q3          Max
## -2.676205354 -0.636885362 -0.005202893  0.586056673  3.342143075
##
## Number of Observations: 332
## Number of Groups: 6
summary(sUAS_ndvi_model)$tTable

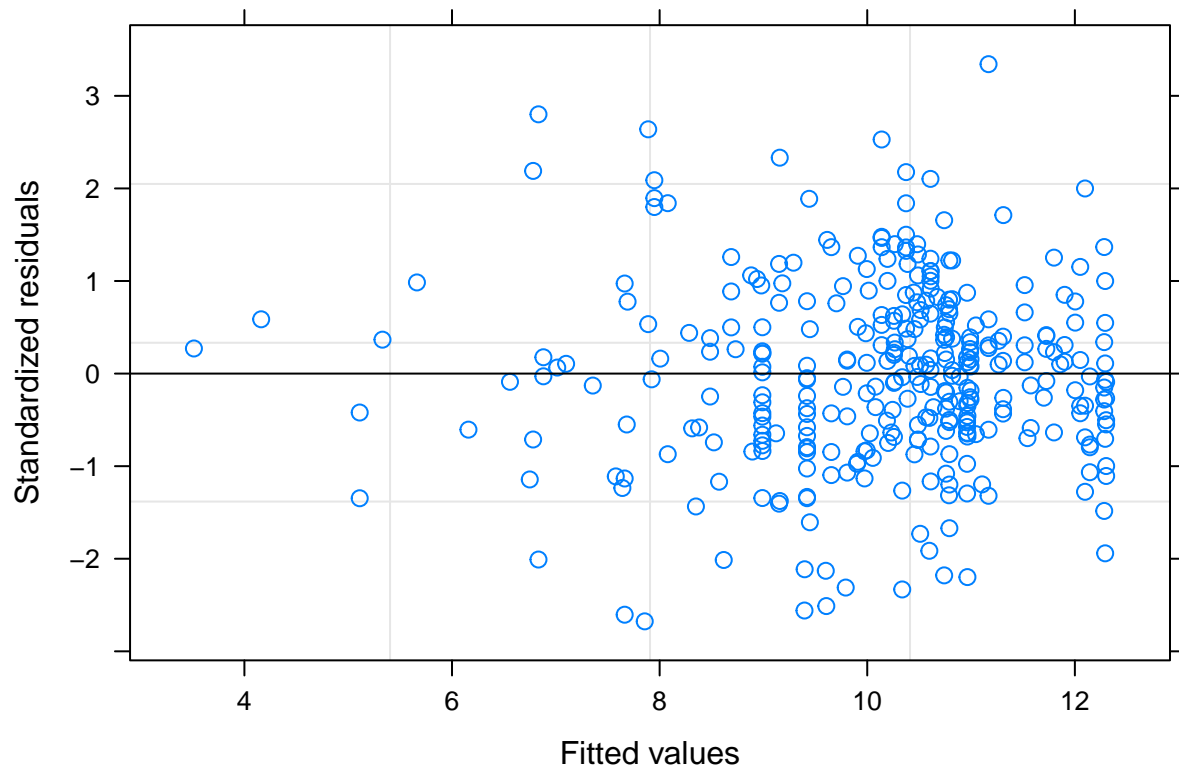
##              Value Std.Error DF   t-value      p-value
## (Intercept)   -24.3921810  7.97649532 322 -3.058007 2.414970e-03
## SI             47.6344646 17.70619487 322  2.690271 7.511739e-03
## TopDress_kgha   0.2085855  0.04286881 322  4.865671 1.788676e-06
## I(SI * SI)     -12.5387959  9.81534736 322 -1.277468 2.023572e-01
## SI:TopDress_kgha -0.2026276  0.04450175 322 -4.553251 7.501114e-06
Anova(sUAS_ndvi_model , type = 3)

## Analysis of Deviance Table (Type III tests)
##
## Response: GrainYield_Mgha
##              Chisq Df Pr(>Chisq)
## (Intercept)    9.3514 1  0.002228 **
## SI              7.2376 1  0.007139 **
## TopDress_kgha  23.6748 1 1.141e-06 ***
## I(SI * SI)      1.6319 1  0.201437
## SI:TopDress_kgha 20.7321 1 5.282e-06 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
r.squaredGLMM(sUAS_ndvi_model)
```

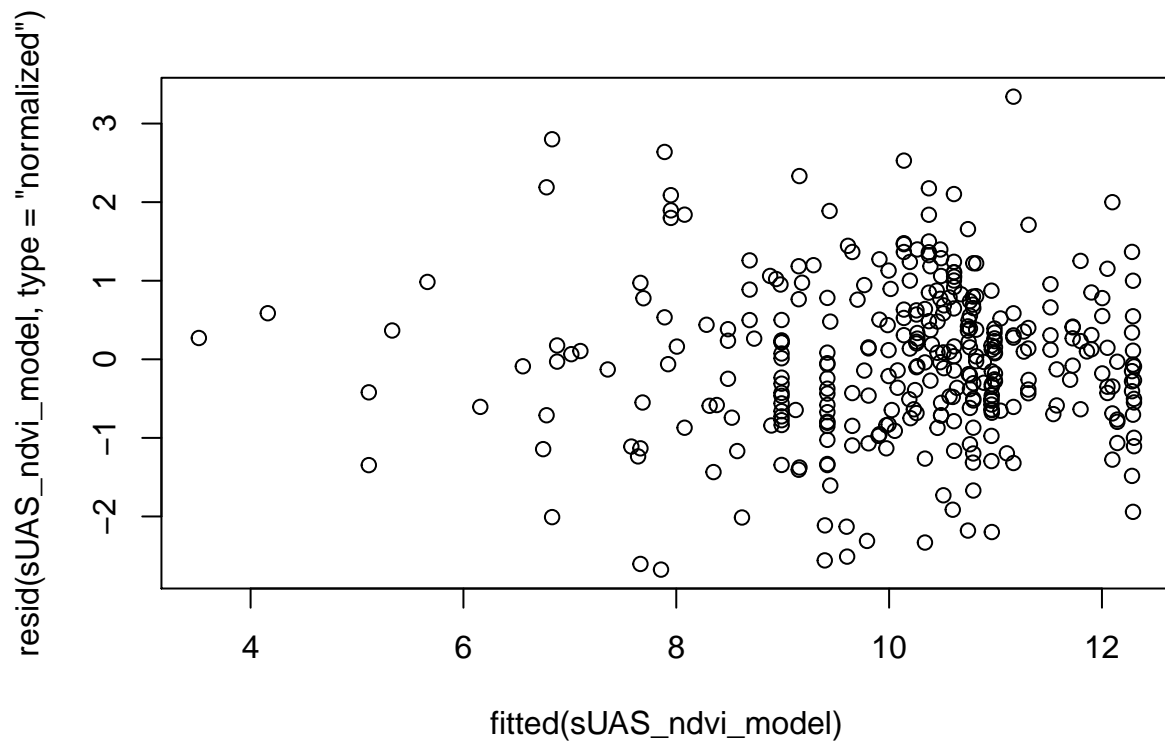
```
##           R2m      R2c
## [1,] 0.471899 0.7985167
```

model diagnostics

```
plot (sUAS_ndvi_model)
```



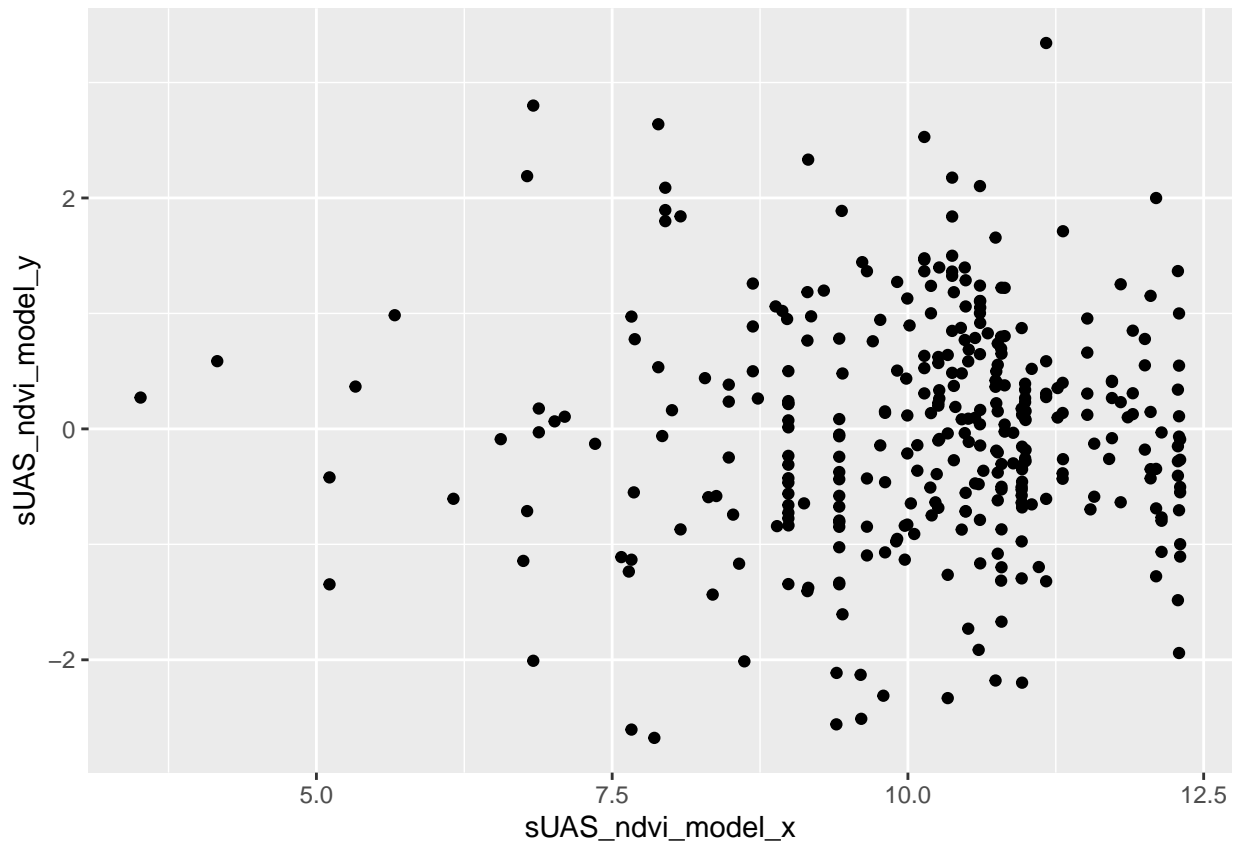
```
plot(resid(sUAS_ndvi_model, type = "normalized") ~fitted(sUAS_ndvi_model))
```



```
sUAS_ndvi_model_y <- resid(sUAS_ndvi_model, type = "normalized")
sUAS_ndvi_model_x <- fitted(sUAS_ndvi_model)

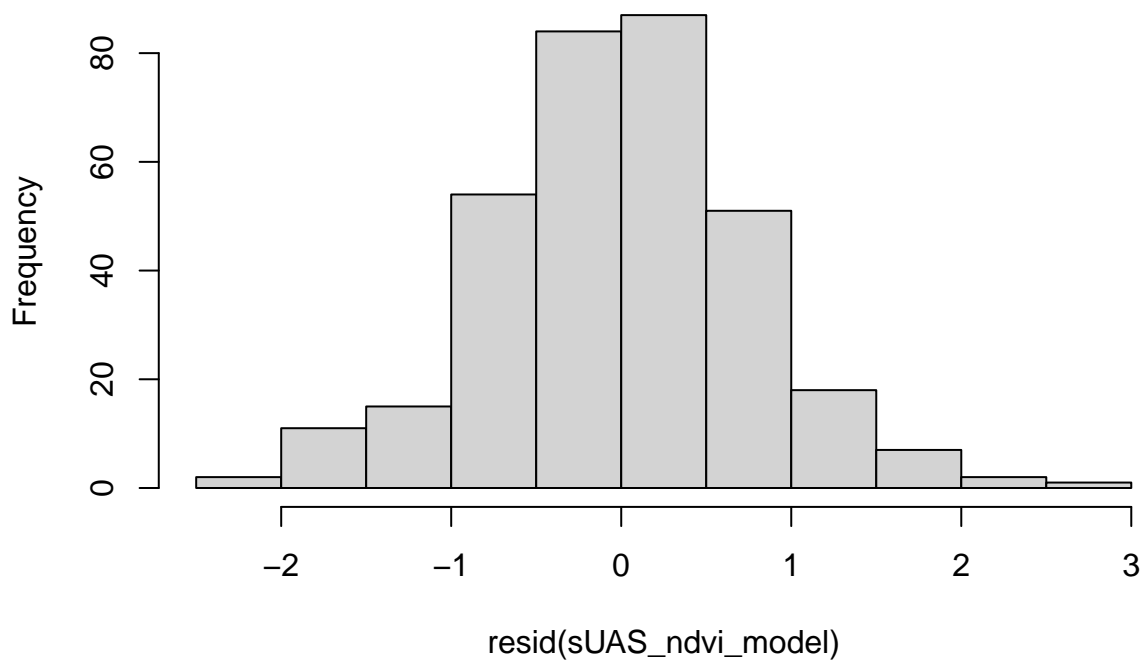
sUAS_ndvi_modelresid_data <- data.frame(sUAS_ndvi_model_x , sUAS_ndvi_model_y)

ggplot( data = sUAS_ndvi_modelresid_data , aes( x = sUAS_ndvi_model_x , y = sUAS_ndvi_model_y)) +
  geom_point(mapping = aes(sUAS_ndvi_model_x , sUAS_ndvi_model_y) , data = sUAS_ndvi_modelresid_data)
```

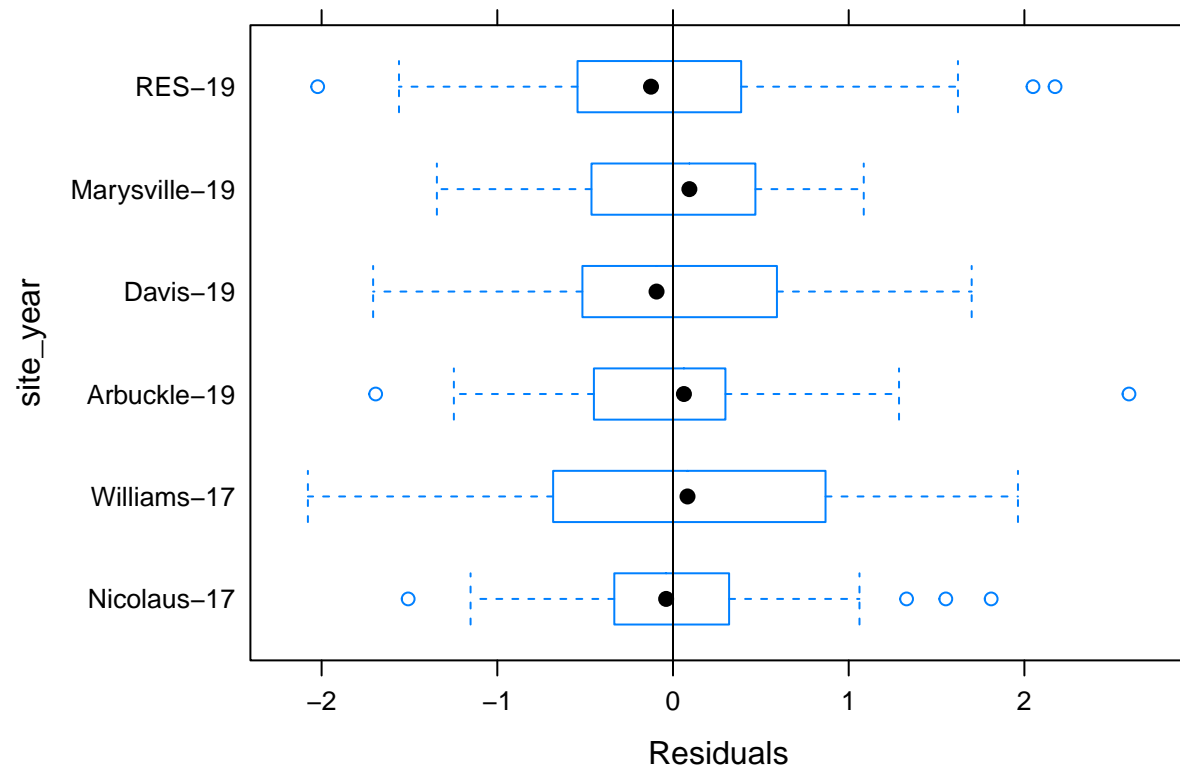


```
hist(resid(sUAS_ndvi_model))
```

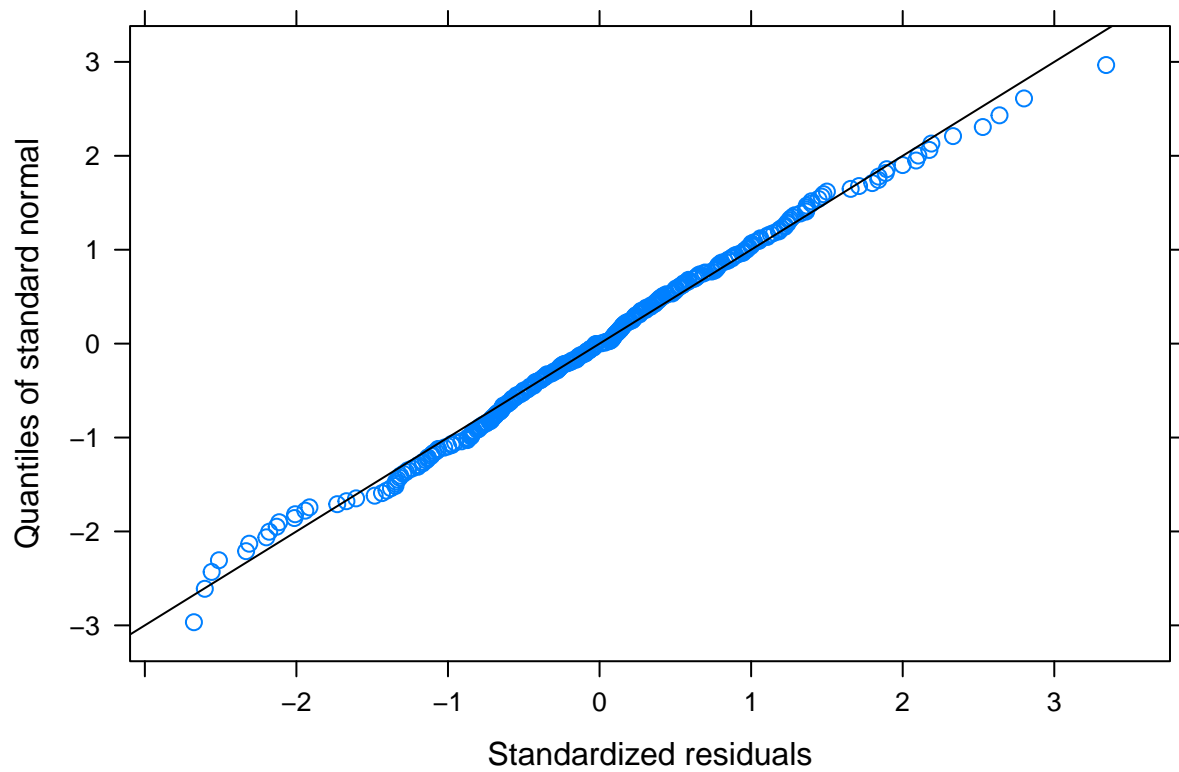
Histogram of resid(sUAS_ndvi_model)



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

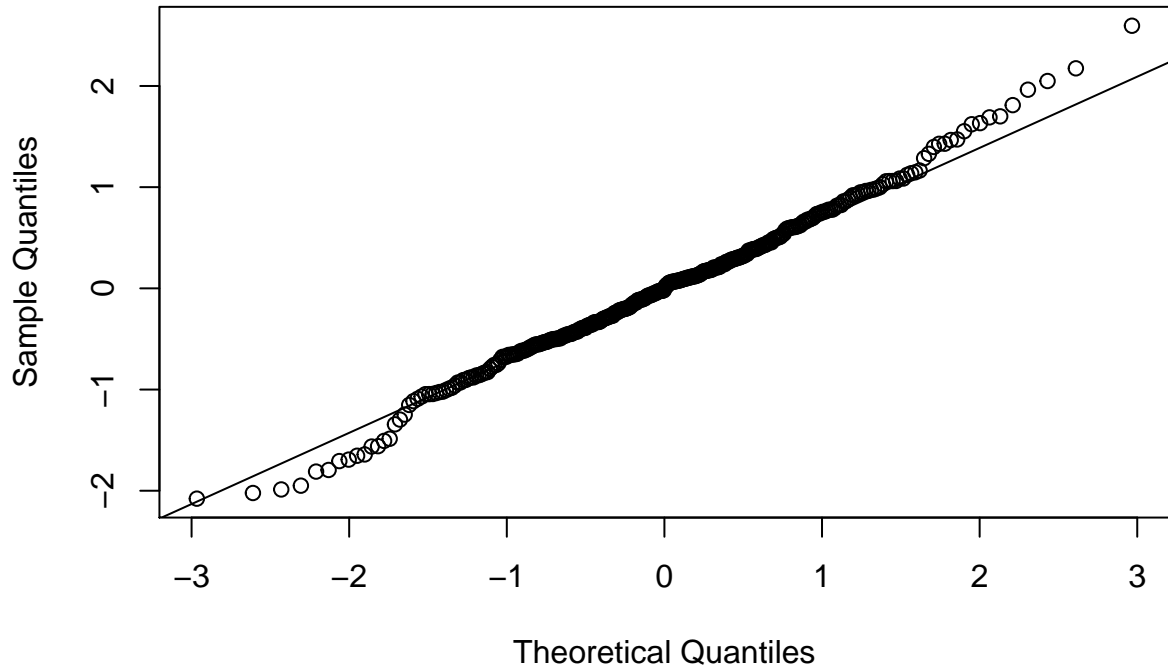


```
qqnorm(sUAS_ndvi_model, abline = c(0,1) )
```

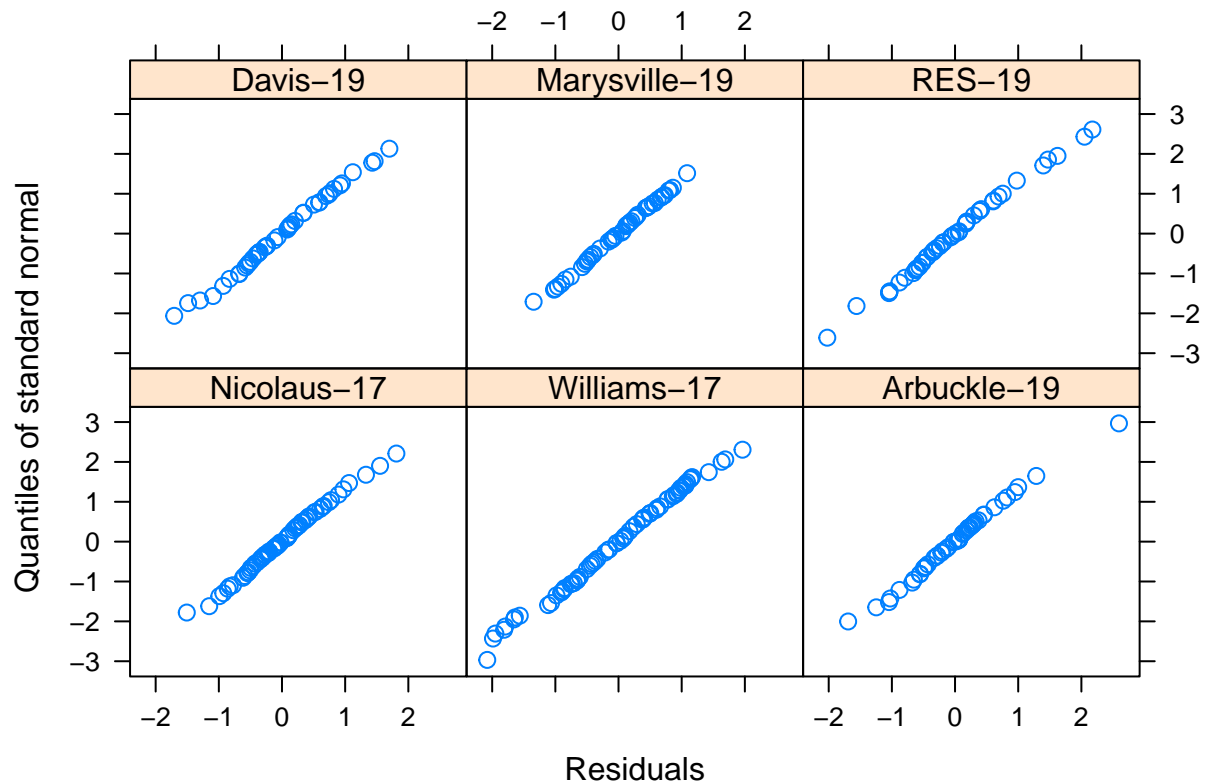



```
qqnorm(resid(sUAS_ndvi_model))
qqline(resid(sUAS_ndvi_model))
```

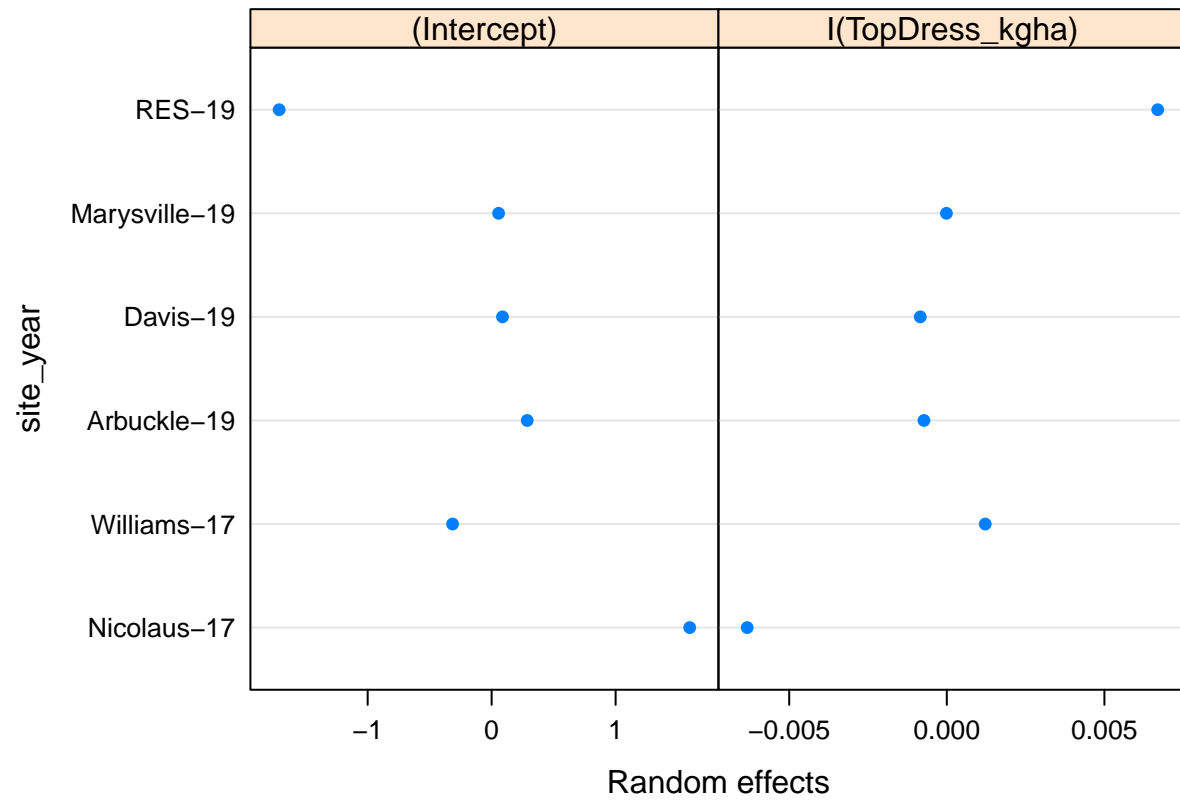
Normal Q-Q Plot



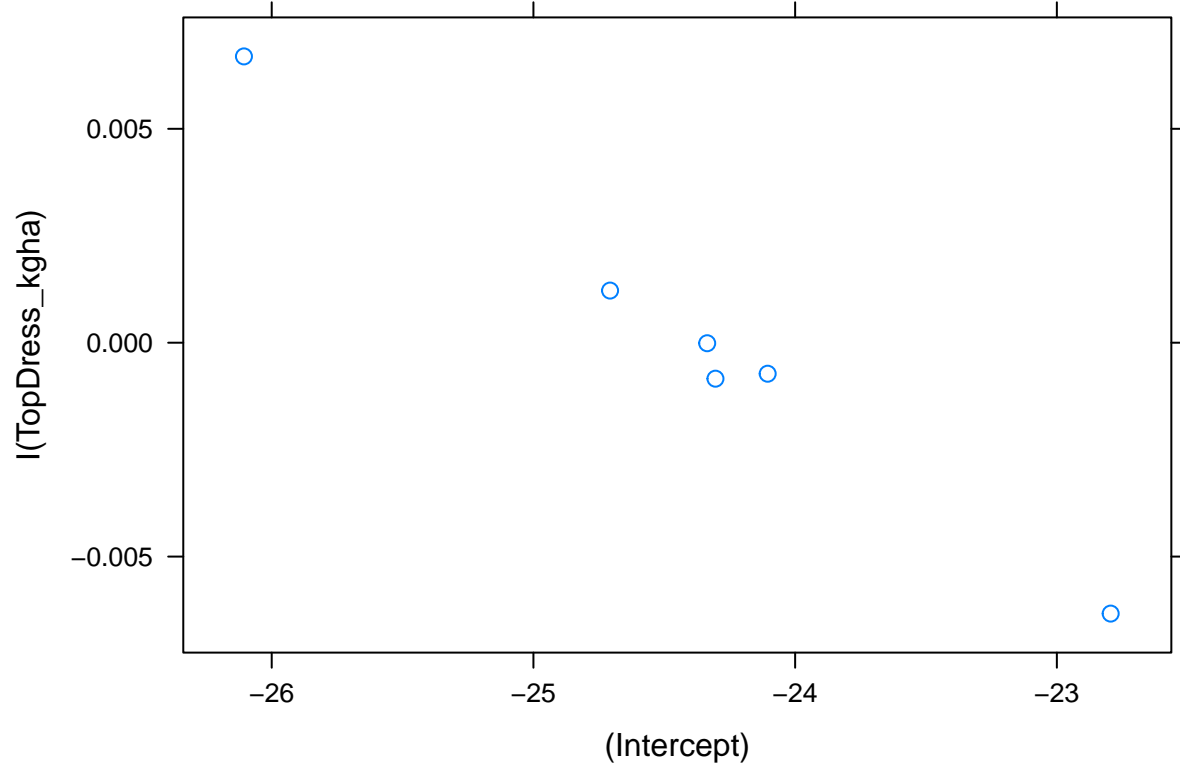
```
qqnorm(sUAS_ndvi_model , ~resid(.) | site_year)
```



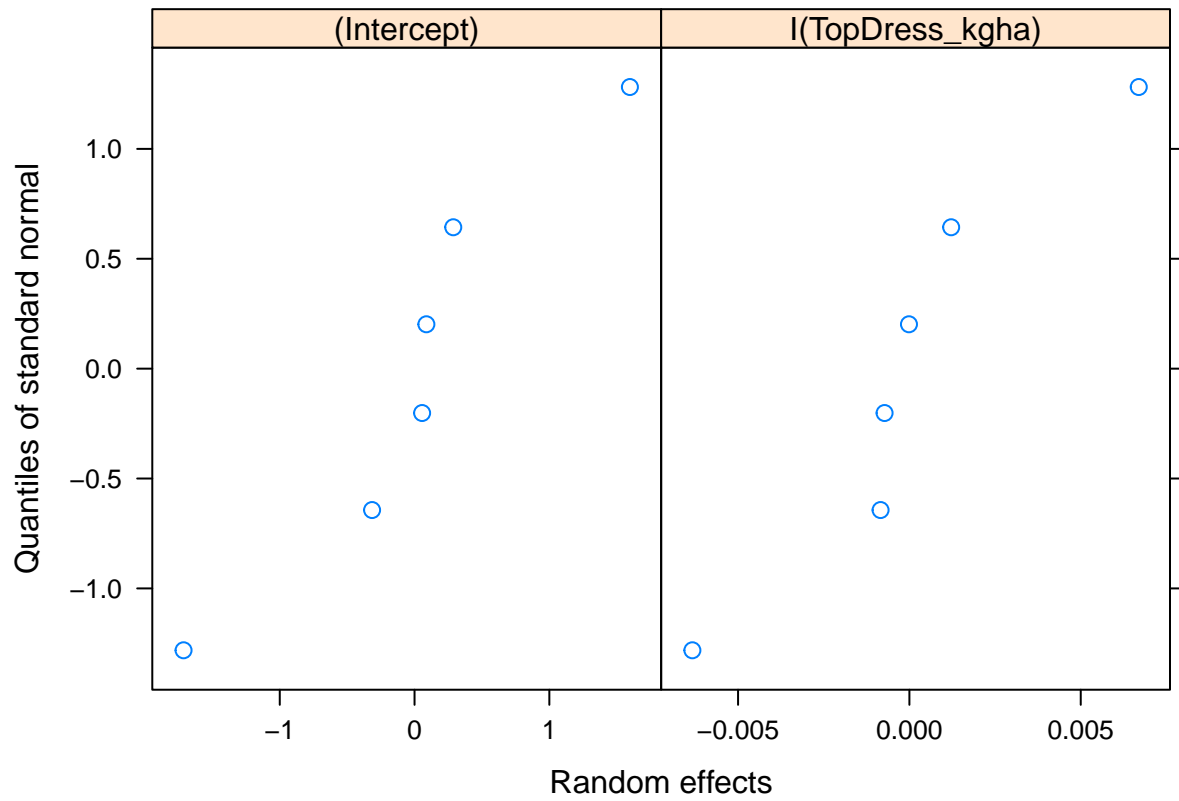
```
plot(ranef(sUAS_ndvi_model))
```



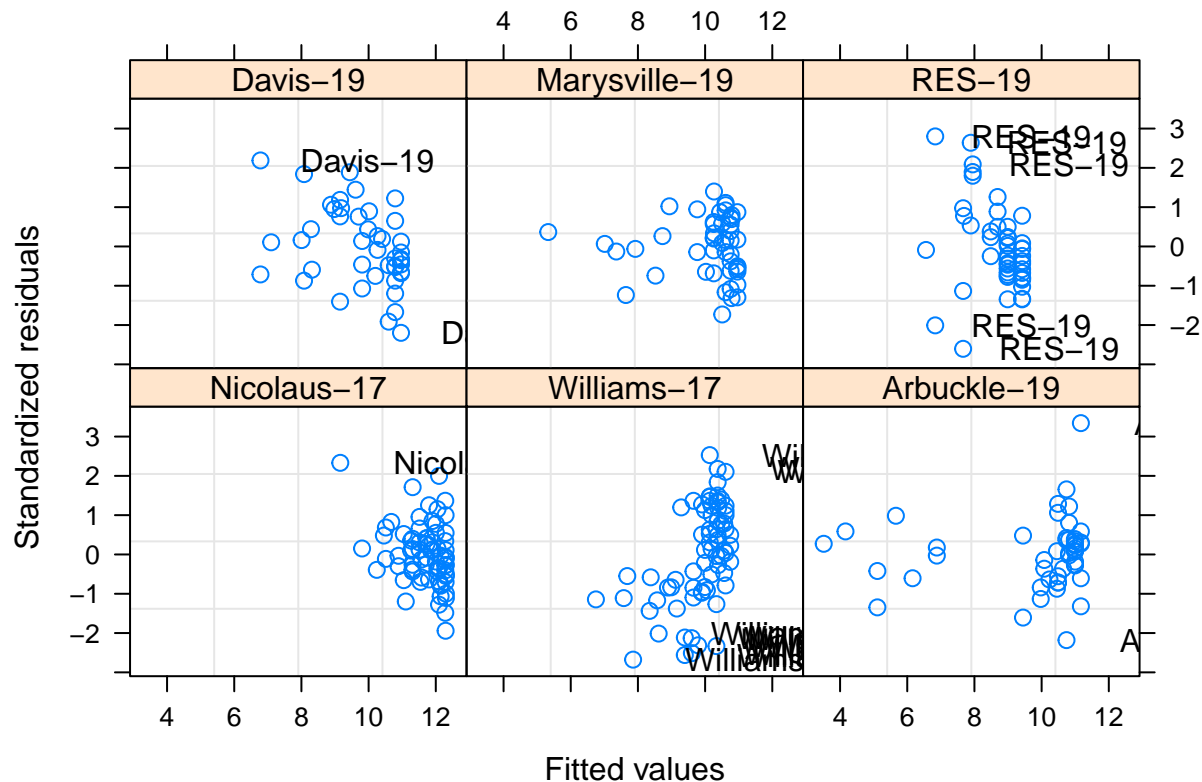
```
pairs(sUAS_ndvi_model , id = 0.1)
```



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



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



```
###outlier removal
```

```
sUAS_ndvi_data_OR <- sUAS_ndvi_data[-(c(which(abs(residuals(sUAS_ndvi_model,type="normalized"))>qnorm(0
```

```
###rerun model sans outliers
```

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

```
sUAS_ndvi_model_OR <- lme(GrainYield_Mgha ~ SI * TopDress_kgha + I(SI*SI) ,
  control = ctrl ,
  random = ~ I(TopDress_kgha) | site_year ,
  data = sUAS_ndvi_data)
```

```
summary(sUAS_ndvi_model_OR)
```

```
## Linear mixed-effects model fit by REML
```

```
## Data: sUAS_ndvi_data
```

```
## AIC BIC logLik
```

```
## 821.447 855.5566 -401.7235
```

```
##
```

```
## Random effects:
```

```
## Formula: ~I(TopDress_kgha) | site_year
```

```
## Structure: General positive-definite, Log-Cholesky parametrization
```

```
## StdDev Corr
```

```
## (Intercept) 1.071047781 (Intr)
```

```
## I(TopDress_kgha) 0.004528693 -0.935
```

```
## Residual 0.776505570
```

```
##
```

```
## Fixed effects: GrainYield_Mgha ~ SI * TopDress_kgha + I(SI * SI)
```

```
## Value Std.Error DF t-value p-value
```

```
## (Intercept)      -24.39218  7.976495 322 -3.058007  0.0024
## SI                47.63446 17.706195 322  2.690271  0.0075
## TopDress_kgha     0.20859  0.042869 322  4.865671  0.0000
## I(SI * SI)       -12.53880  9.815347 322 -1.277468  0.2024
## SI:TopDress_kgha -0.20263  0.044502 322 -4.553251  0.0000
## Correlation:
##              (Intr) SI      TpDrs_ I(SI*S
## SI              -0.996
## TopDress_kgha   -0.025 -0.031
## I(SI * SI)       0.989 -0.998  0.078
## SI:TopDress_kgha 0.023  0.031 -0.998 -0.078
##
## Standardized Within-Group Residuals:
##              Min          Q1          Med          Q3          Max
## -2.676205354 -0.636885362 -0.005202893  0.586056673  3.342143075
##
## Number of Observations: 332
## Number of Groups: 6
```

```
summary(sUAS_ndvi_model_OR)$tTable
```

```
##              Value Std.Error DF   t-value    p-value
## (Intercept)  -24.3921810  7.97649532 322 -3.058007 2.414970e-03
## SI            47.6344646 17.70619487 322  2.690271 7.511739e-03
## TopDress_kgha  0.2085855  0.04286881 322  4.865671 1.788676e-06
## I(SI * SI)    -12.5387959  9.81534736 322 -1.277468 2.023572e-01
## SI:TopDress_kgha -0.2026276  0.04450175 322 -4.553251 7.501114e-06
```

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

```
## Analysis of Deviance Table (Type III tests)
```

```
##
```

```
## Response: GrainYield_Mgha
```

```
##              Chisq Df Pr(>Chisq)
## (Intercept)    9.3514 1  0.002228 **
## SI              7.2376 1  0.007139 **
## TopDress_kgha  23.6748 1  1.141e-06 ***
## I(SI * SI)      1.6319 1  0.201437
## SI:TopDress_kgha 20.7321 1  5.282e-06 ***
```

```
## ---
```

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

```
r.squaredGLMM(sUAS_ndvi_model_OR)
```

```
##              R2m      R2c
```

```
## [1,] 0.471899 0.7985167
```

```
ndvi_r_sq <- r.squaredGLMM(sUAS_ndvi_model_OR)
```

```
ndvi_r_sq_fixed <- round(ndvi_r_sq[1] , digits = 2)
```

```
ndvi_r_sq_fixed
```

```
## [1] 0.47
```

```
ndvi_r_sq_total <- round(ndvi_r_sq[2] , digits = 2)
```

```
ndvi_r_sq_total
```

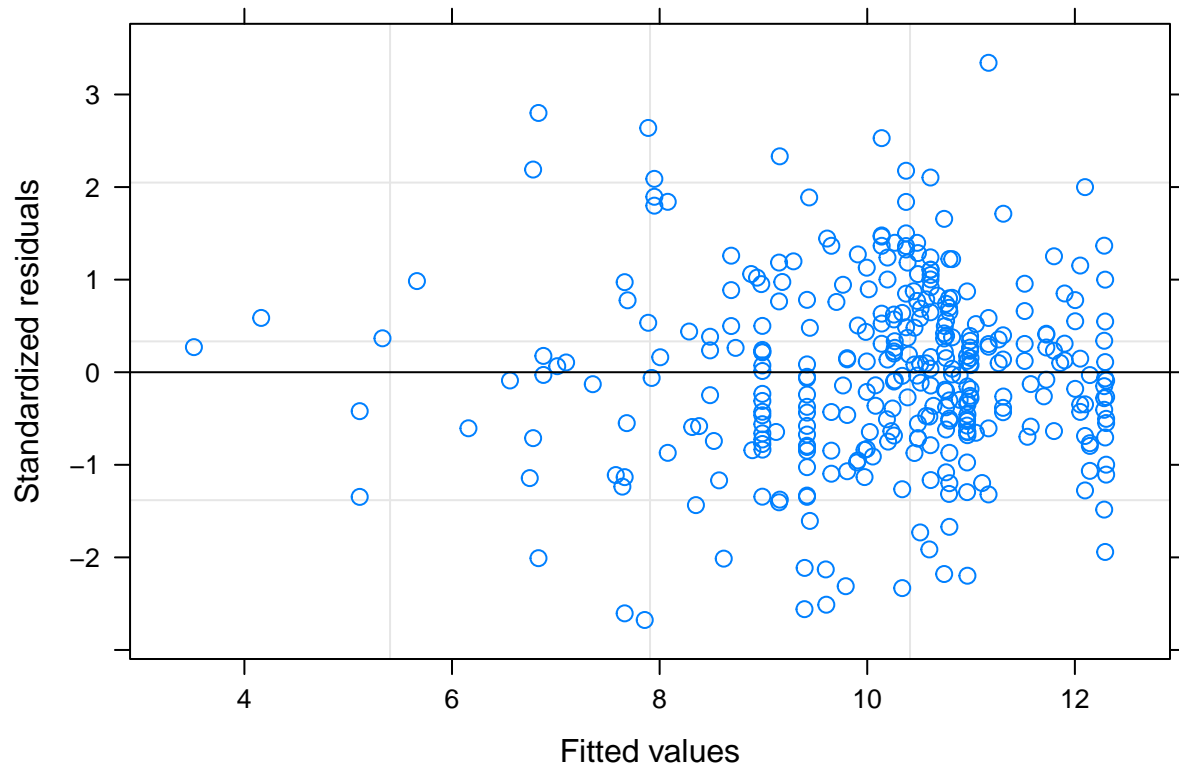
```
## [1] 0.8
```

```
ndvi_r_sq_random <- ndvi_r_sq_total - ndvi_r_sq_fixed  
ndvi_r_sq_random
```

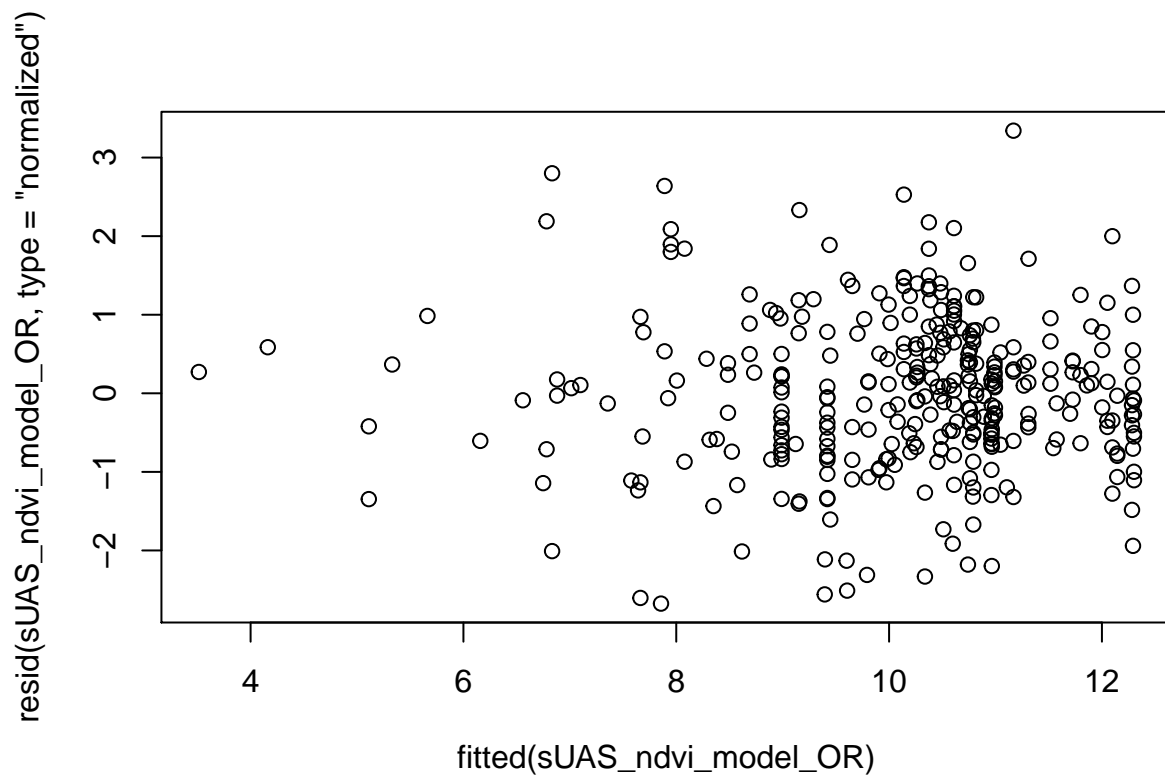
```
## [1] 0.33
```

model diagnostics

```
plot (sUAS_ndvi_model_OR)
```



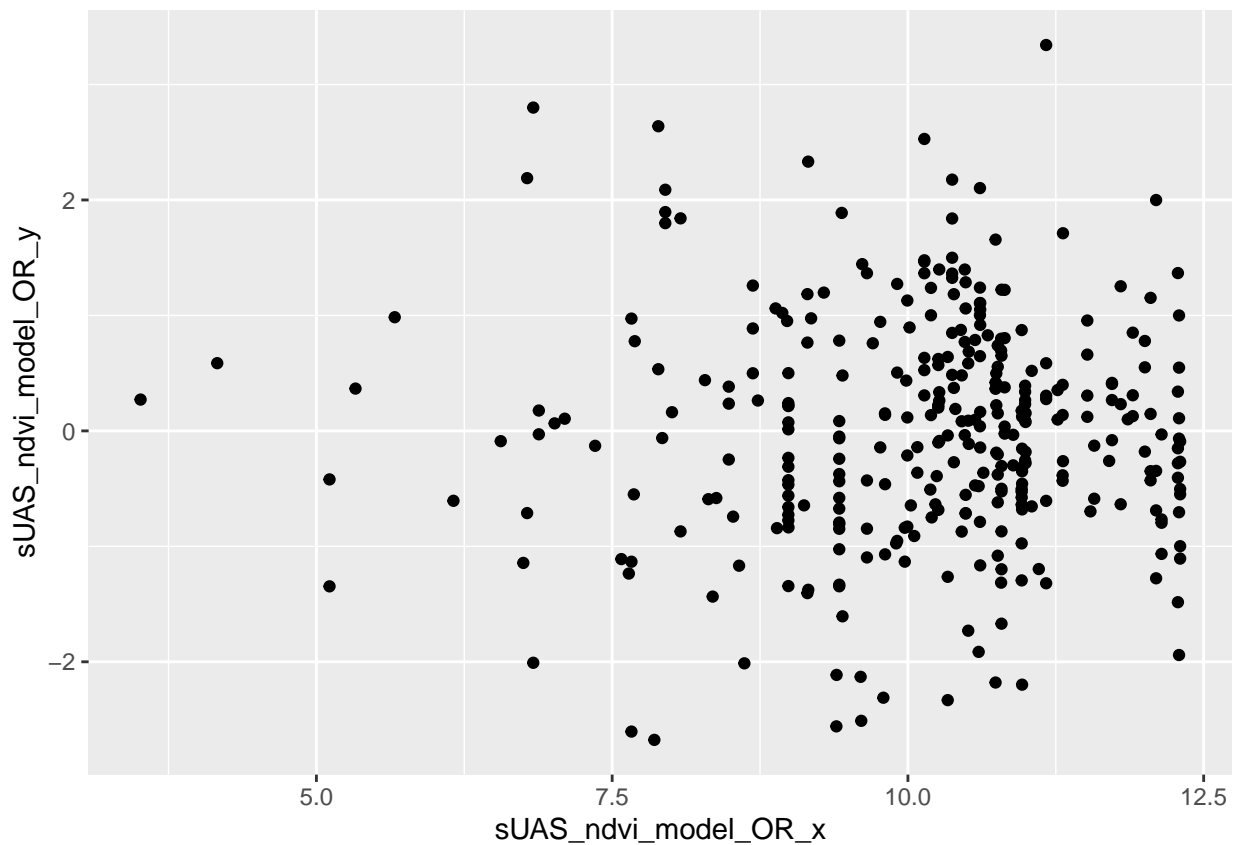
```
plot(resid(sUAS_ndvi_model_OR, type = "normalized") ~fitted(sUAS_ndvi_model_OR))
```



```
sUAS_ndvi_model_OR_y <- resid(sUAS_ndvi_model_OR, type = "normalized")
sUAS_ndvi_model_OR_x <- fitted(sUAS_ndvi_model_OR)

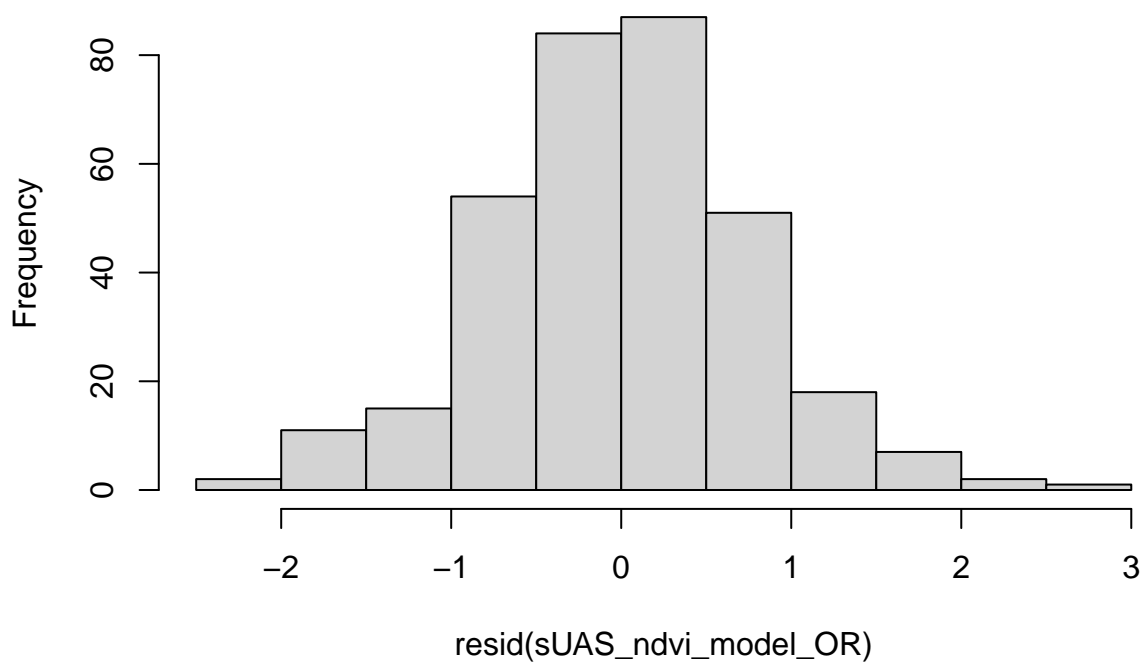
sUAS_ndvi_model_ORresid_data <- data.frame(sUAS_ndvi_model_OR_x , sUAS_ndvi_model_OR_y)

ggplot( data = sUAS_ndvi_model_ORresid_data , aes( x = sUAS_ndvi_model_OR_x , y = sUAS_ndvi_model_OR_y))
  geom_point(mapping = aes(sUAS_ndvi_model_OR_x , sUAS_ndvi_model_OR_y) , data = sUAS_ndvi_model_ORresid_data)
```

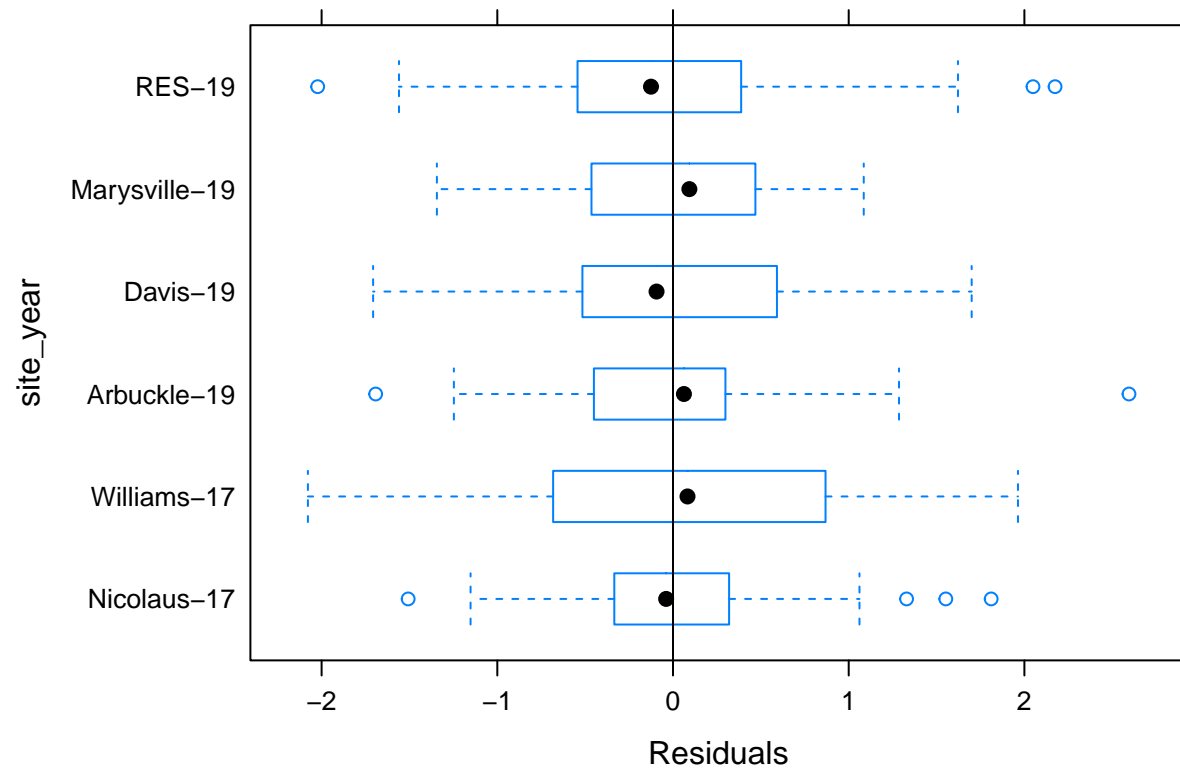


```
hist(resid(sUAS_ndvi_model_OR))
```

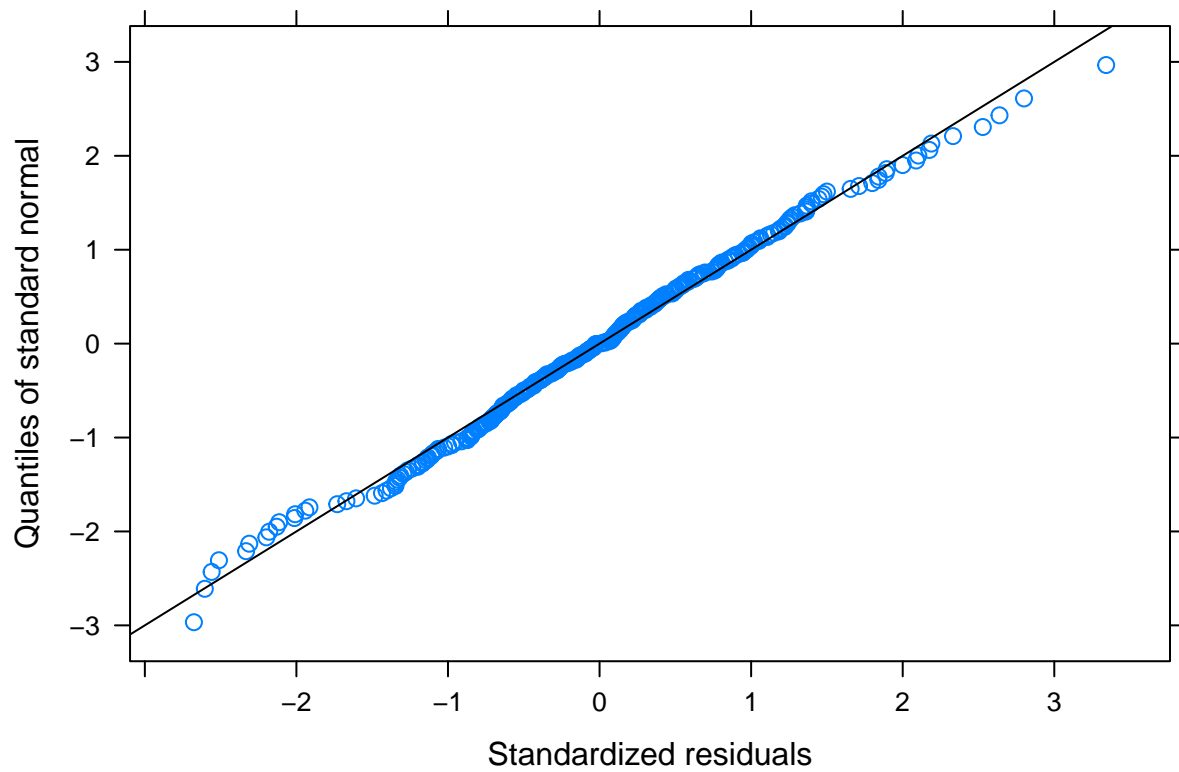
Histogram of resid(sUAS_ndvi_model_OR)




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

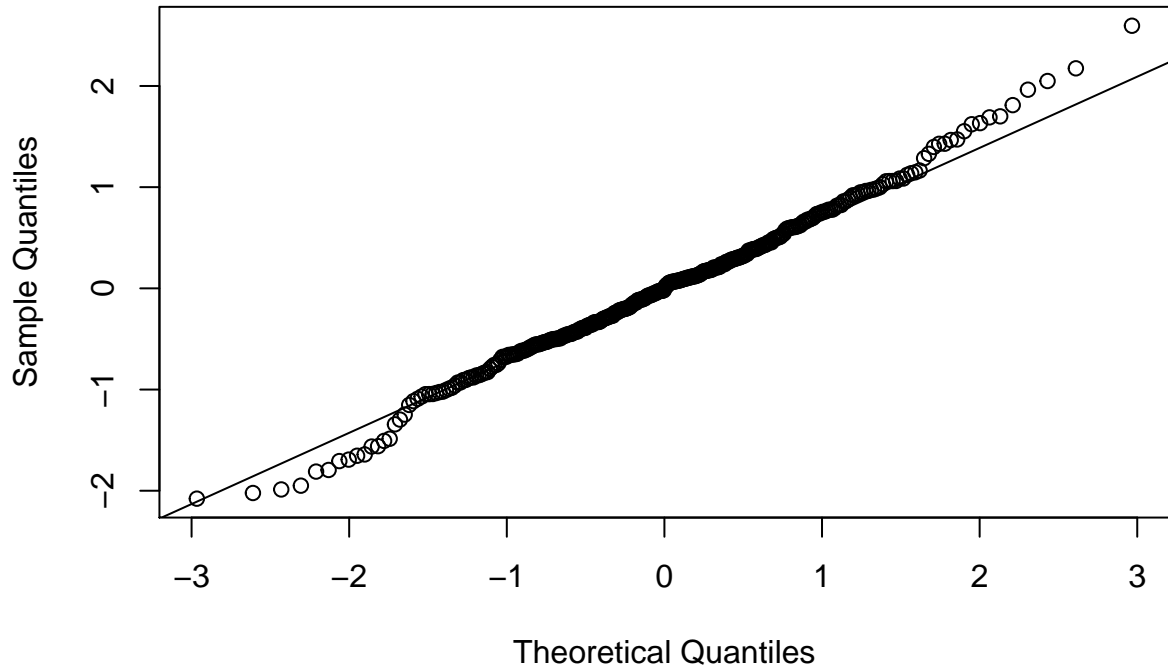


```
qqnorm(sUAS_ndvi_model_OR, abline = c(0,1) )
```

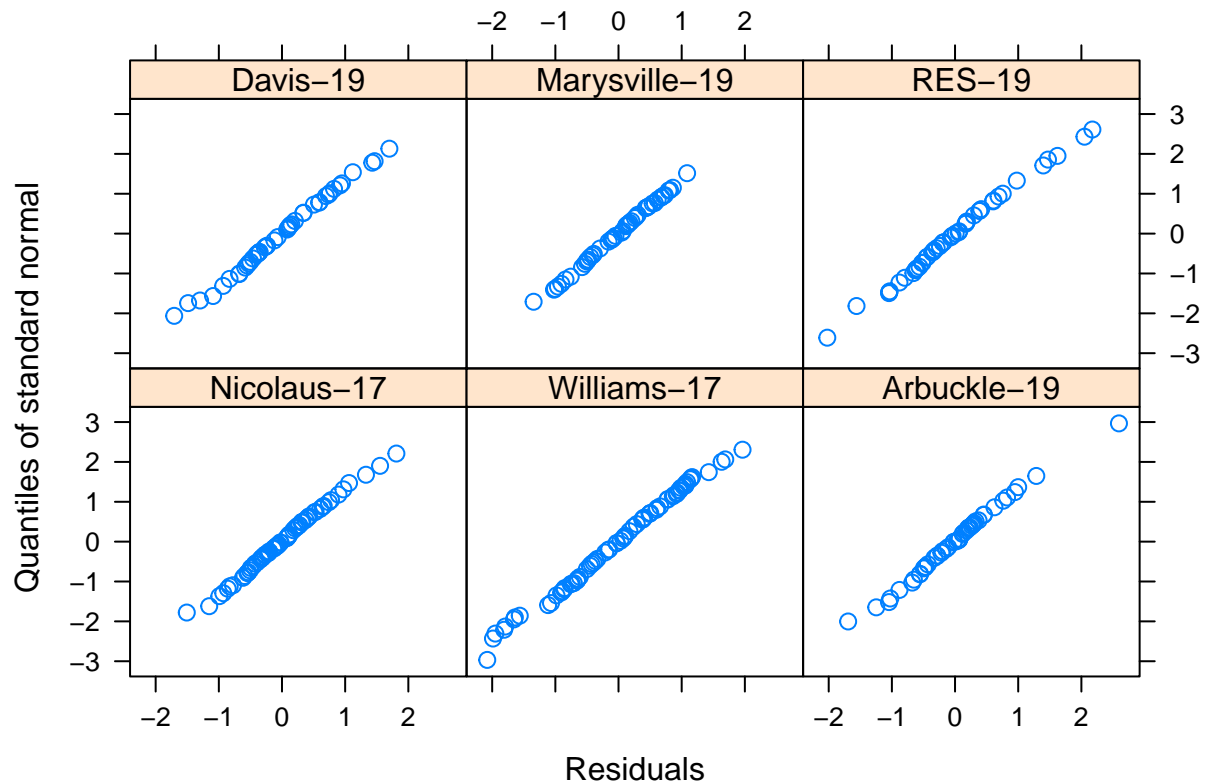


```
qqnorm(resid(sUAS_ndvi_model_OR))
qqline(resid(sUAS_ndvi_model_OR))
```

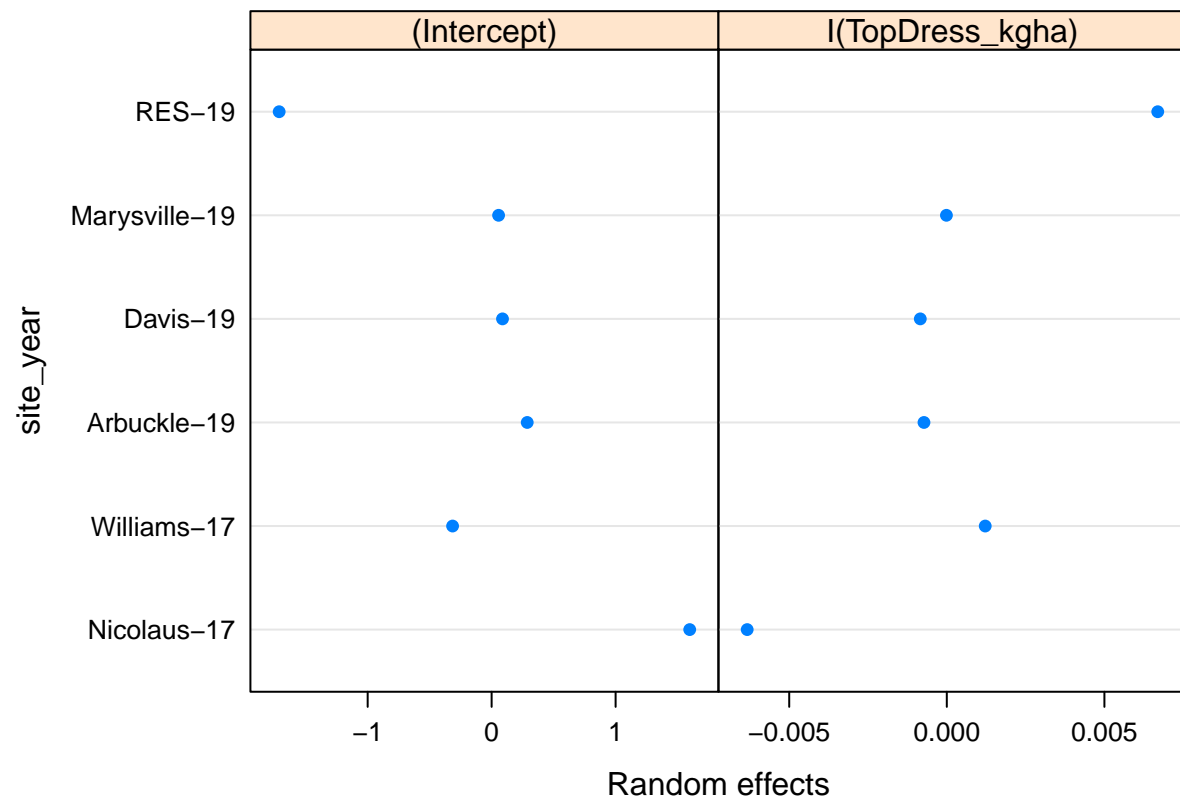
Normal Q-Q Plot



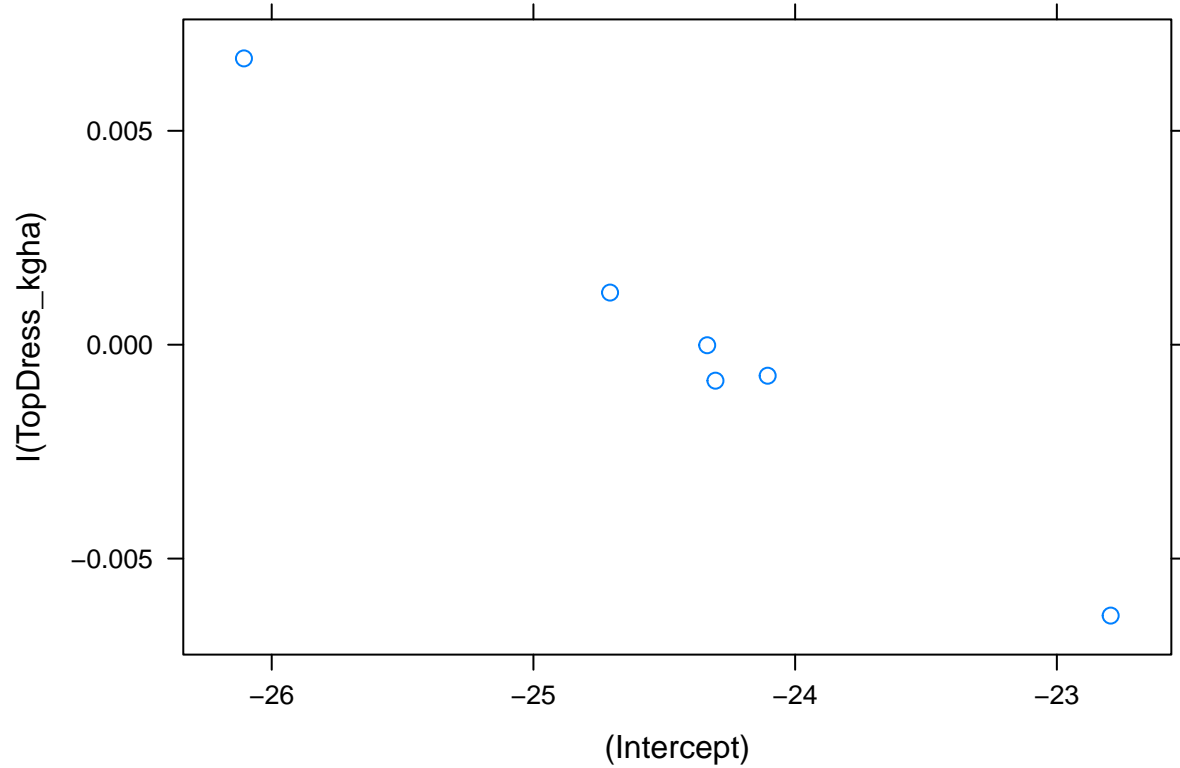
```
qqnorm(sUAS_ndvi_model_OR , ~resid(.) | site_year)
```



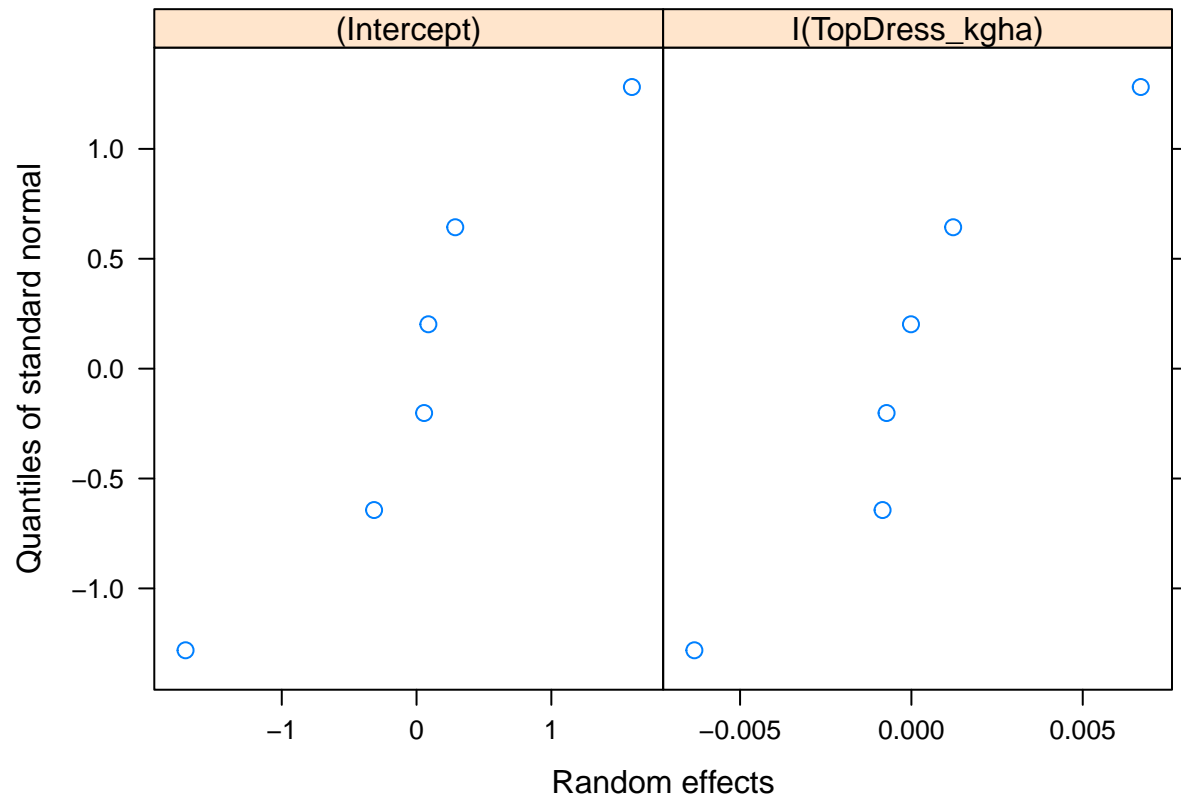
```
plot(ranef(sUAS_ndvi_model_OR))
```



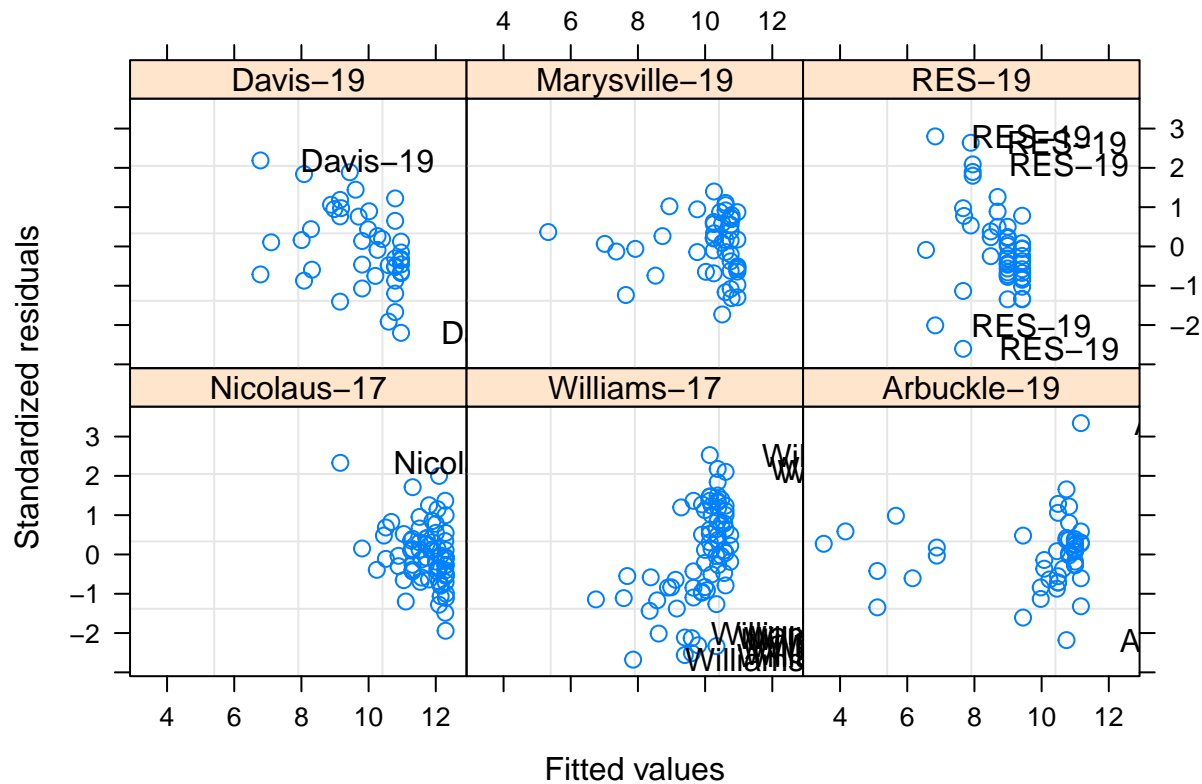
```
pairs(sUAS_ndvi_model_OR , id = 0.1)
```



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



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



```
### emmeans
```

```
mylist <- list(SI=seq(0.708 , 1 , by = .0001), TopDress_kgha = c( 34 , 0))
```

```
sUAS_ndvi_emmeans <- emmeans(sUAS_ndvi_model_OR , ~TopDress_kgha * SI , at = mylist )
```

```
sUAS_ndvi_emmeans_contrast <- as.data.frame(summary(contrast(sUAS_ndvi_emmeans , "pairwise" , side = ">"))
```

```
sUAS_ndvi_emmeans_contrast <- sUAS_ndvi_emmeans_contrast %>%  
  mutate(sUAS_ndvi_Response_Index = 1 / SI )
```

```
sUAS_ndvi_emmeans_contrast <- sUAS_ndvi_emmeans_contrast %>%  
  mutate(t_score = abs((estimate - 0.26)) / SE)
```

```
sUAS_ndvi_emmeans_contrast <- sUAS_ndvi_emmeans_contrast %>%  
  mutate(prob_greater_than_26 = if_else(estimate < 0.26 , pt(q = t_score , df = df , lower.tail = F) , 1 - pt(q = t_score , df = df , lower.tail = F))
```

```
sUAS_ndvi_emmeans_contrast <- sUAS_ndvi_emmeans_contrast %>%  
  mutate(prob_diff_from_26 = 1 - (pt(q = t_score , df = df , lower.tail = F) + pt(q = -t_score , df = df , lower.tail = F))
```

```
sUAS_ndvi_emmeans_contrast$sUAS_ndvi_Response_Index_r <- (round(sUAS_ndvi_emmeans_contrast$sUAS_ndvi_Response_Index_r , 2))
```

```
sUAS_ndvi_run1 <- 125 - 100
```

```
sUAS_ndvi_rise1 <- 1.596946 - 0.2038182
```

```
sUAS_ndvi_slope1 <- round((sUAS_ndvi_rise1 / sUAS_ndvi_run1) * 5 , digits = 2)
```

```
sUAS_ndvi_slope1
```

```
## [1] 0.28
```

```

sUAS_ndvi_run2 <- 142 - 125
sUAS_ndvi_rise2 <- 2.237784 - 1.596946
sUAS_ndvi_slope2 <- round((sUAS_ndvi_rise2 / sUAS_ndvi_run2) * 5 , digits = 3)
sUAS_ndvi_slope2

## [1] 0.188

sUAS_ndvi_mean_se <- sUAS_ndvi_emmeans_contrast %>%
  select(SE) %>%
  summarise(mean_se = mean(SE))

sUAS_ndvi_mean_se$mean_se <- round(sUAS_ndvi_mean_se$mean_se , digits = 2)

sUAS_ndvi_mean_se$mean_se

## [1] 0.21

###confidence interval

#at every 0.10 interval, what is the 90% confidence interval of the yield response, and what RI value d

str(sUAS_ndvi_emmeans_contrast , give.attr = F)

## 'data.frame': 2921 obs. of 12 variables:
## $ contrast : Factor w/ 1 level "34 - 0": 1 1 1 1 1 1 1 1 1 1 ...
## $ SI : num 0.708 0.708 0.708 0.708 0.708 ...
## $ estimate : num 2.21 2.21 2.21 2.21 2.21 ...
## $ SE : num 0.396 0.396 0.395 0.395 0.395 ...
## $ df : num 322 322 322 322 322 322 322 322 322 322 ...
## $ t.ratio : num 5.59 5.6 5.6 5.6 5.6 ...
## $ p.value : num 2.36e-08 2.36e-08 2.35e-08 2.35e-08 2.35e-08 ...
## $ sUAS_ndvi_Response_Index : num 1.41 1.41 1.41 1.41 1.41 ...
## $ t_score : num 4.94 4.94 4.94 4.94 4.94 ...
## $ prob_greater_than_26 : num 1 1 1 1 1 ...
## $ prob_diff_from_26 : num 1 1 1 1 1 ...
## $ sUAS_ndvi_Response_Index_r: num 1.41 1.41 1.41 1.41 1.41 ...

sUAS_ndvi_emmeans_contrast <- sUAS_ndvi_emmeans_contrast %>%
  mutate(response_mgha_r = round(estimate , digits = 2),
         se_mgha_r = round(SE , digits = 3))

#estimate = 0.20 , SE = 0.114

print(round(0.20 - z_score*0.114 , digits = 2))

## [1] 0.01

print(round(0.20 + z_score*0.114 , digits = 2))

## [1] 0.39

#estimate = 0.30 , SE = 0.104

print(round(0.30 - z_score*0.106 , digits = 2))

## [1] 0.13

```

```

print(round(0.30 + z_score*0.106 , digits = 2))

## [1] 0.47
#estimate = 0.40 , SE = 0.100

print(round(0.40 - z_score*0.100 , digits = 2))

## [1] 0.24
print(round(0.40 + z_score*0.100 , digits = 2))

## [1] 0.56
#estimate = 0.50 , SE = 0.099

print(round(0.50 - z_score*0.099 , digits = 2))

## [1] 0.34
print(round(0.50 + z_score*0.099 , digits = 2))

## [1] 0.66
##sUAS NDRE-RI
sUAS_ndre_data <- paper3_data_no_2018 %>%
  filter(Platform == "sUAS_NDRE")

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

sUAS_ndre_model <- lme(GrainYield_Mgha ~ SI * TopDress_kgha + I(SI*SI) ,
  control = ctrl ,
  random = ~ I(TopDress_kgha) | site_year ,
  data = sUAS_ndre_data)

summary(sUAS_ndre_model)

## Linear mixed-effects model fit by REML
##   Data: sUAS_ndre_data
##       AIC      BIC    logLik
##  784.3945 818.5041 -383.1972
##
## Random effects:
## Formula: ~I(TopDress_kgha) | site_year
## Structure: General positive-definite, Log-Cholesky parametrization
##              StdDev      Corr
## (Intercept)  1.025351819 (Intr)
## I(TopDress_kgha) 0.003962301 -0.919
## Residual      0.726916252
##
## Fixed effects: GrainYield_Mgha ~ SI * TopDress_kgha + I(SI * SI)
##              Value Std.Error DF   t-value p-value
## (Intercept)  -11.45611  1.717014 322  -6.672109     0
## SI           39.19829  4.123213 322   9.506734     0
## TopDress_kgha  0.10343  0.016790 322   6.160088     0
## I(SI * SI)    -16.99240  2.527712 322  -6.722443     0
## SI:TopDress_kgha -0.10031  0.018537 322  -5.411511     0

```

```
## Correlation:
##          (Intr) SI      TpDrs_ I(SI*S
## SI          -0.958
## TopDress_kgha -0.138  0.011
## I(SI * SI)    0.934 -0.993  0.075
## SI:TopDress_kgha 0.115 -0.011 -0.987 -0.076
##
## Standardized Within-Group Residuals:
##          Min          Q1          Med          Q3          Max
## -2.607665461 -0.648795155 -0.006631185  0.600700113  3.701360082
##
## Number of Observations: 332
## Number of Groups: 6
```

```
summary(sUAS_ndre_model)$tTable
```

	Value	Std.Error	DF	t-value	p-value
(Intercept)	-11.4561084	1.71701447	322	-6.672109	1.101242e-10
SI	39.1982873	4.12321292	322	9.506734	4.730027e-19
TopDress_kgha	0.1034275	0.01678994	322	6.160088	2.167408e-09
I(SI * SI)	-16.9924007	2.52771225	322	-6.722443	8.140477e-11
SI:TopDress_kgha	-0.1003116	0.01853671	322	-5.411511	1.222445e-07

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

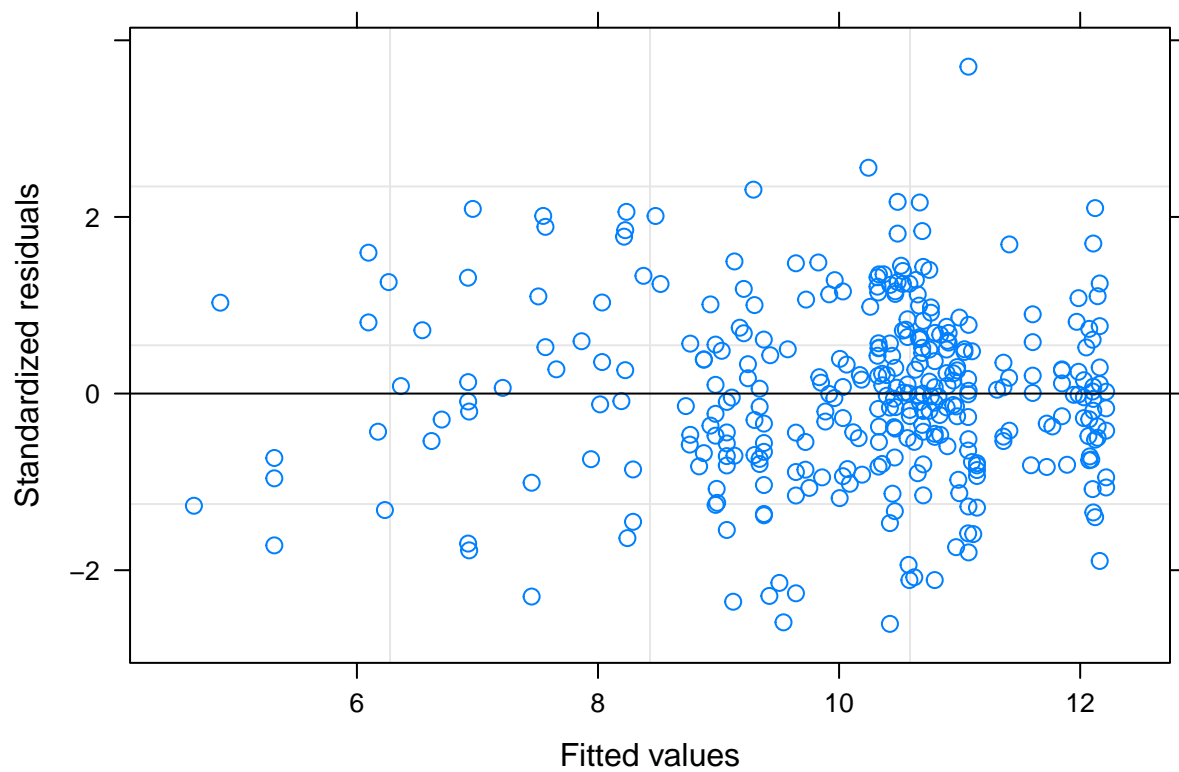
```
## Analysis of Deviance Table (Type III tests)
##
## Response: GrainYield_Mgha
##          Chisq Df Pr(>Chisq)
## (Intercept)  44.517  1 2.522e-11 ***
## SI          90.378  1 < 2.2e-16 ***
## TopDress_kgha 37.947  1 7.270e-10 ***
## I(SI * SI)   45.191  1 1.787e-11 ***
## SI:TopDress_kgha 29.285  1 6.250e-08 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
r.squaredGLMM(sUAS_ndre_model)
```

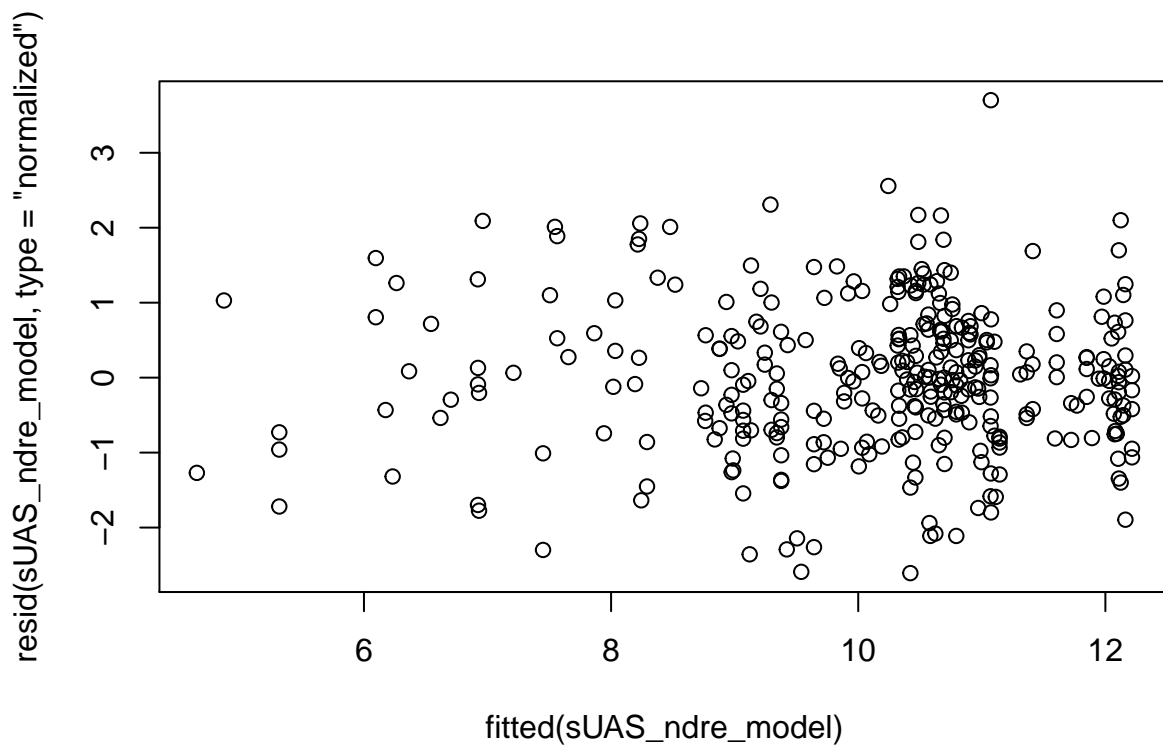
	R2m	R2c
[1,]	0.5079755	0.8193337

model diagnostics

```
plot (sUAS_ndre_model)
```

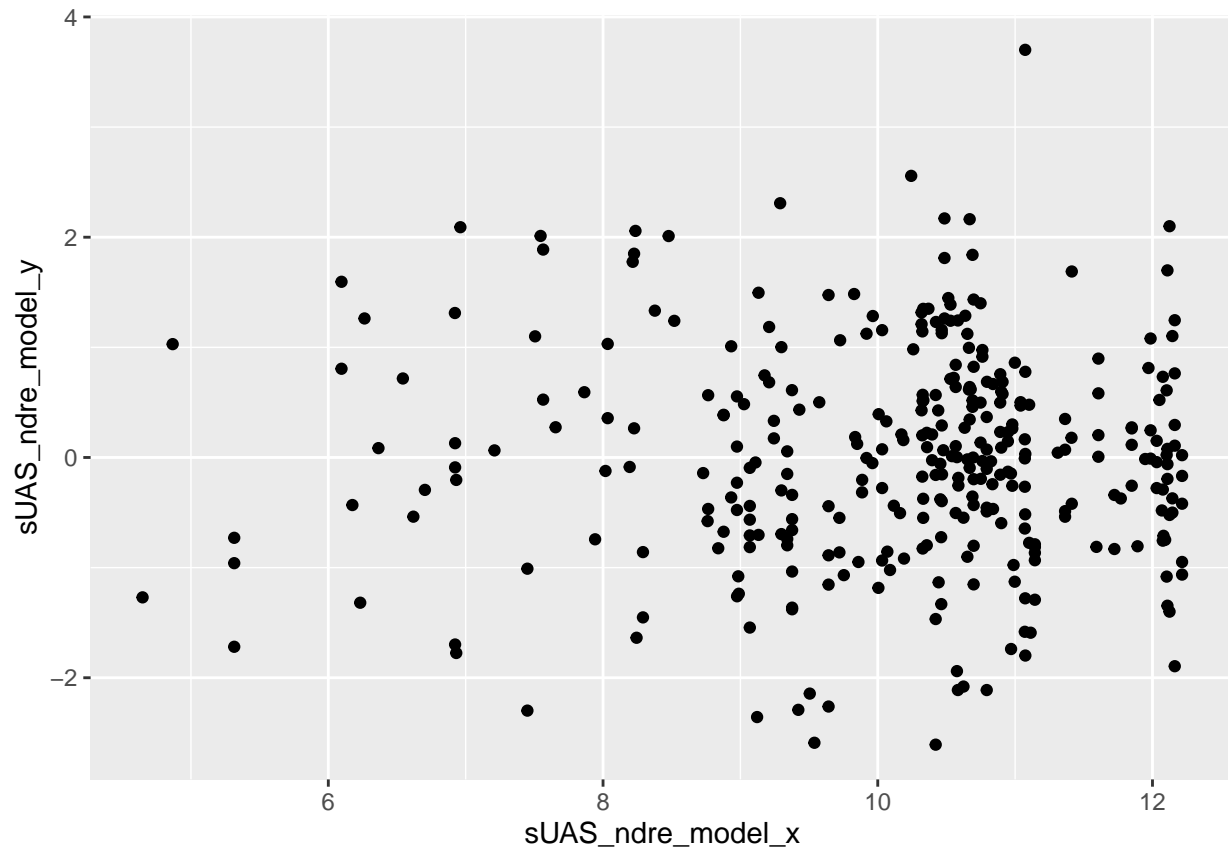
```
plot(resid(sUAS_ndre_model, type = "normalized") ~fitted(sUAS_ndre_model))
```



```
sUAS_ndre_model_y <- resid(sUAS_ndre_model, type = "normalized")
sUAS_ndre_model_x <- fitted(sUAS_ndre_model)

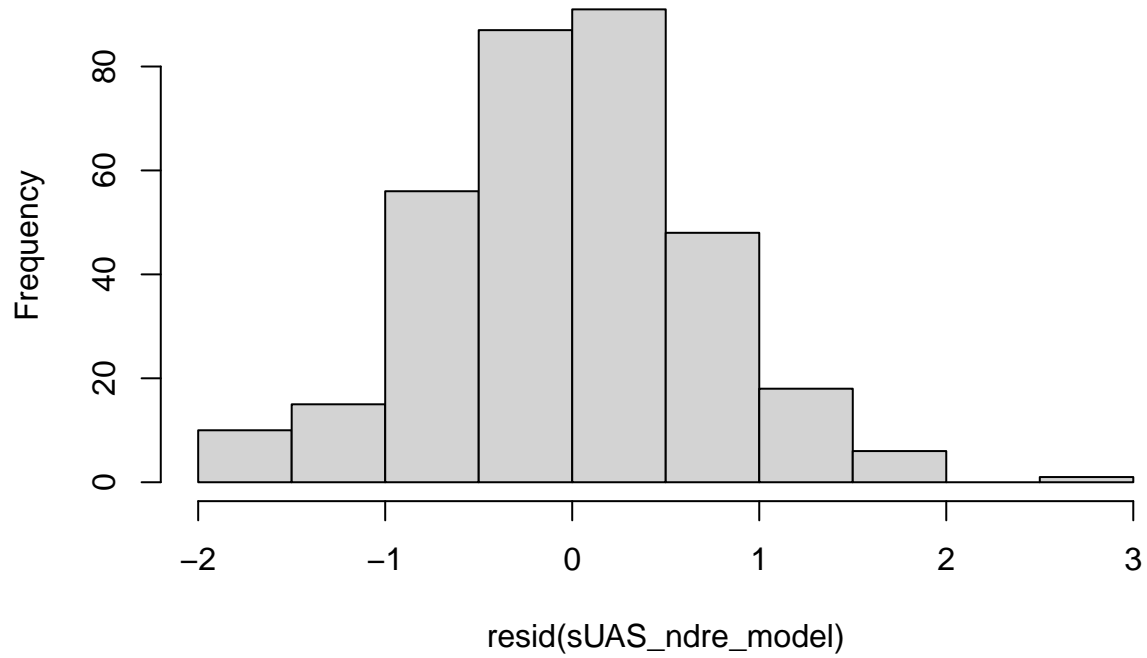
sUAS_ndre_modelresid_data <- data.frame(sUAS_ndre_model_x , sUAS_ndre_model_y)
```

```
ggplot( data = sUAS_ndre_modelresid_data , aes( x = sUAS_ndre_model_x , y = sUAS_ndre_model_y)) +
  geom_point(mapping = aes(sUAS_ndre_model_x , sUAS_ndre_model_y) , data = sUAS_ndre_modelresid_data)
```

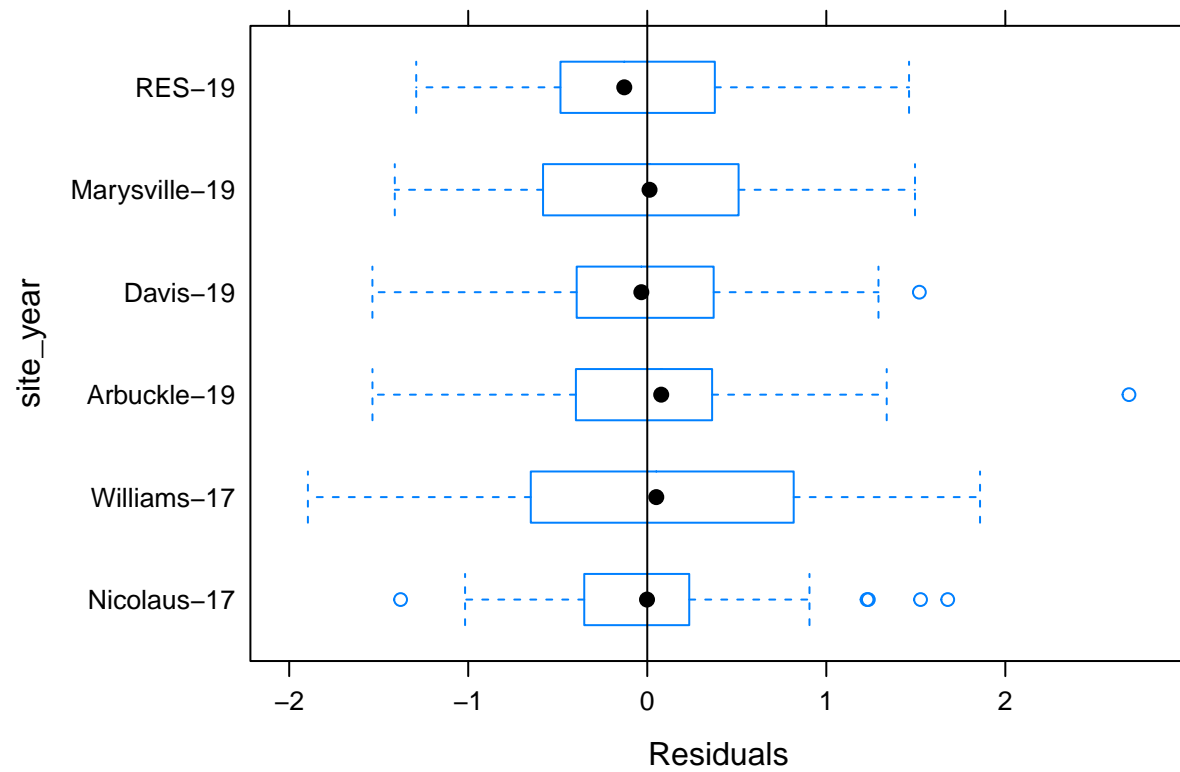


```
hist(resid(sUAS_ndre_model))
```

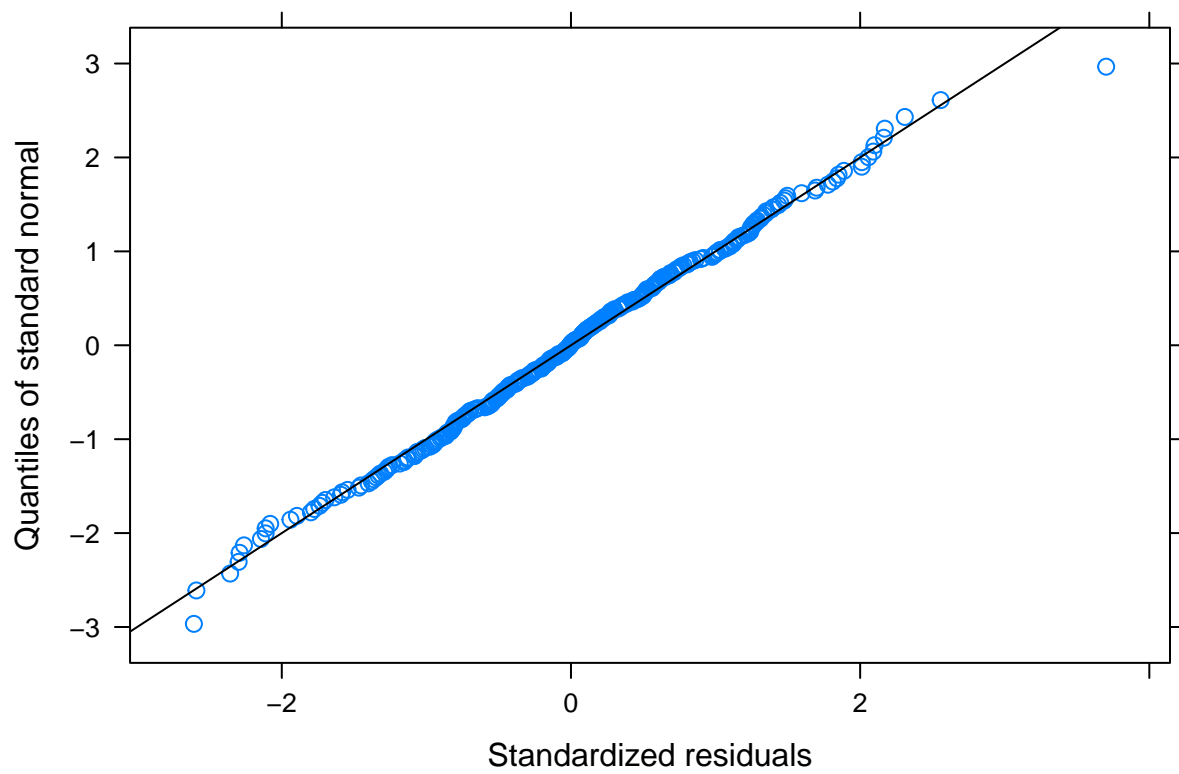
Histogram of resid(sUAS_ndre_model)



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

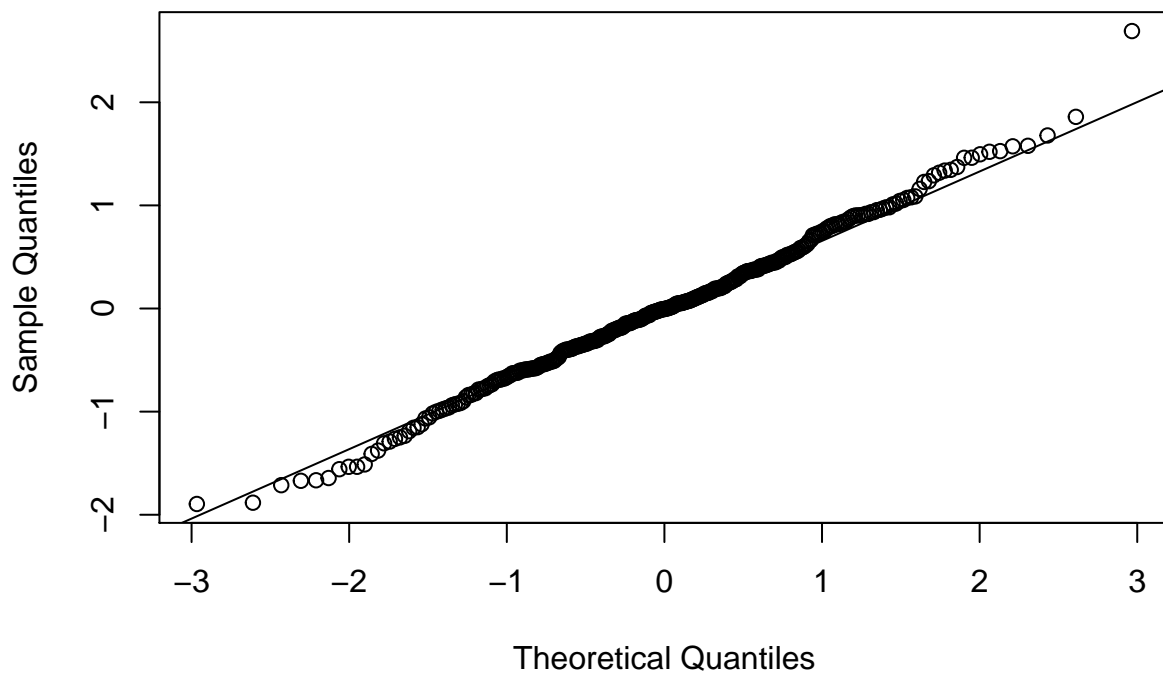


```
qqnorm(sUAS_ndre_model, abline = c(0,1) )
```

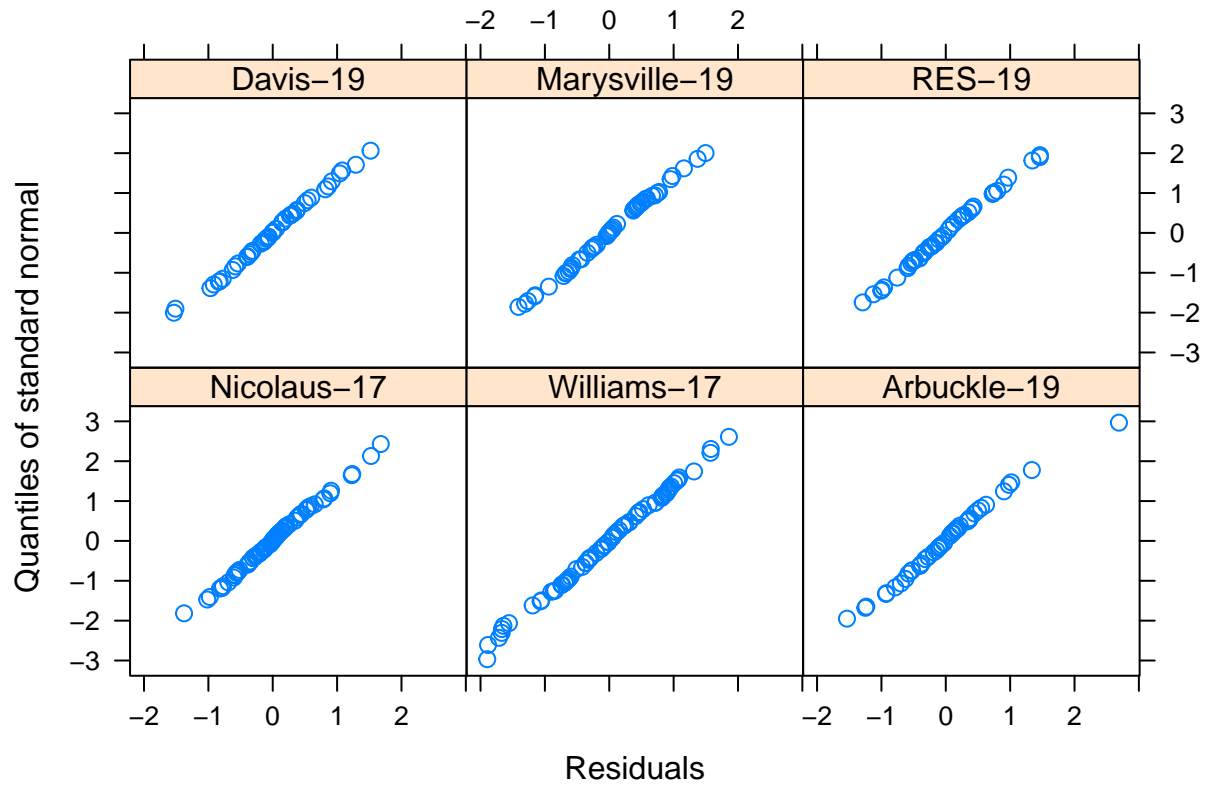


```
qqnorm(resid(sUAS_ndre_model))
qqline(resid(sUAS_ndre_model))
```

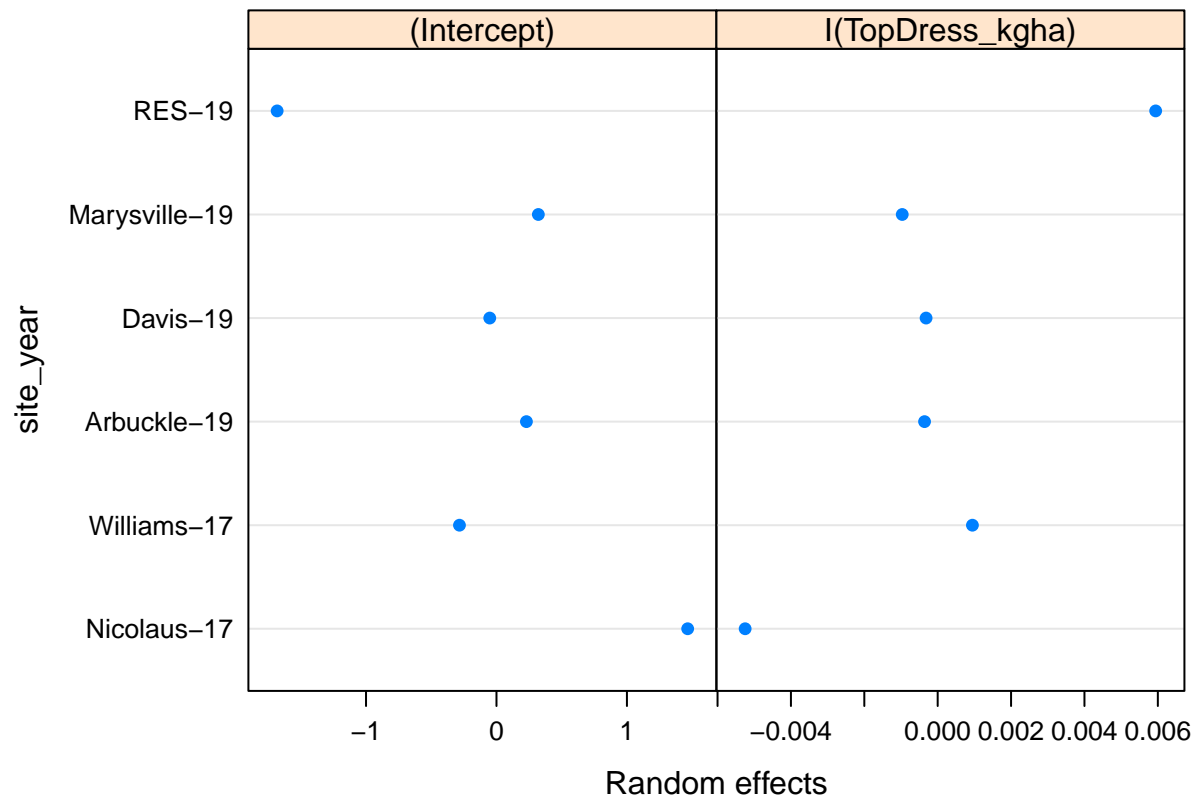
Normal Q–Q Plot



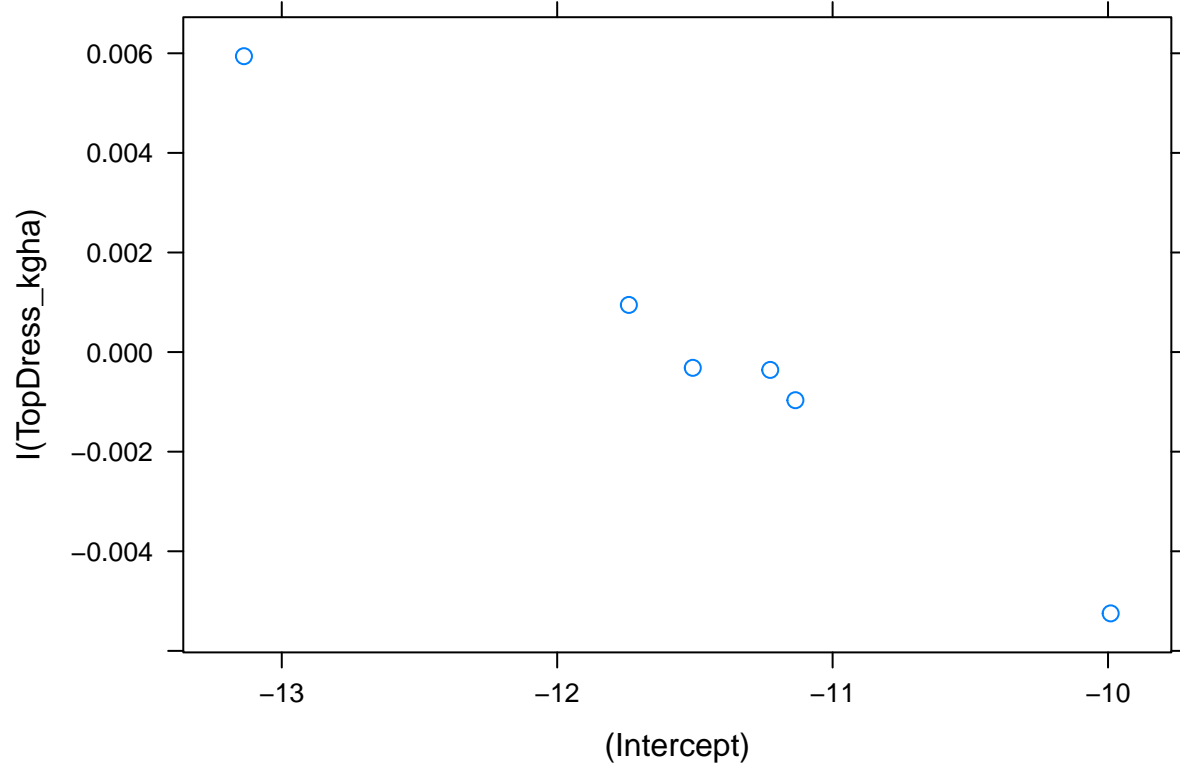
```
qqnorm(sUAS_ndre_model , ~resid(.) | site_year)
```



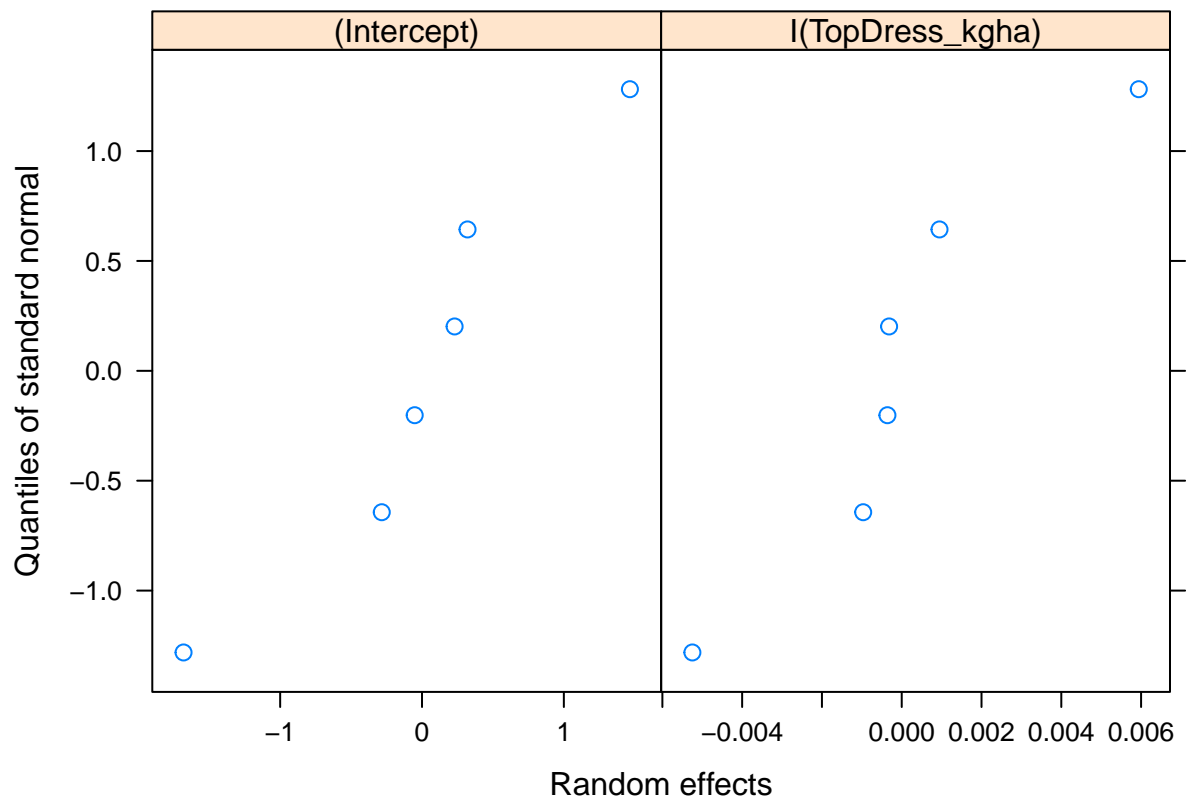
```
plot(ranef(sUAS_ndre_model))
```



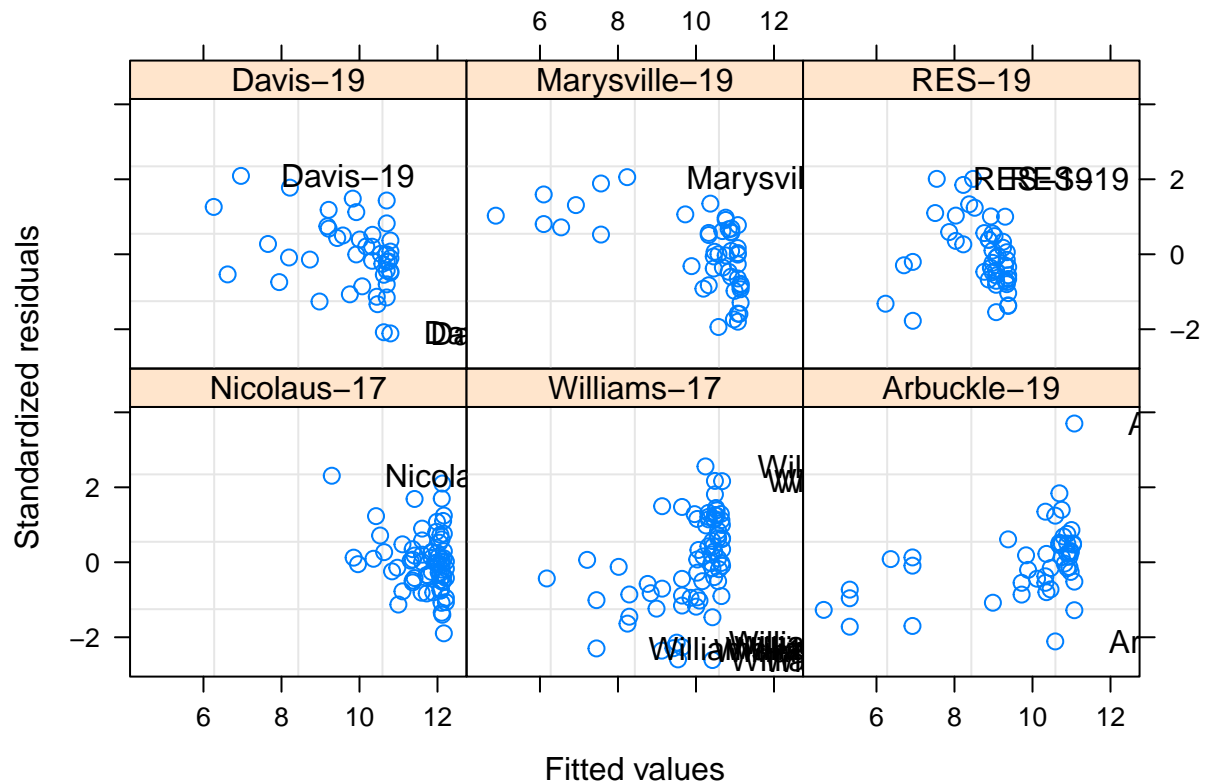
```
pairs(sUAS_ndre_model , id = 0.1)
```



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



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



```
###outlier removal
```

```
sUAS_ndre_data_OR <- sUAS_ndre_data[-(c(which(abs(residuals(sUAS_ndre_model, type = "normalized"))>qnor
```

```
###rerun model sans outliers
```

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

```
sUAS_ndre_model_OR <- lme(GrainYield_Mgha ~ SI * TopDress_kgha + I(SI*SI) ,
      control = ctrl ,
      random = ~ I(TopDress_kgha) | site_year ,
      data = sUAS_ndre_data_OR)
```

```
summary(sUAS_ndre_model_OR)
```

```
## Linear mixed-effects model fit by REML
```

```
## Data: sUAS_ndre_data_OR
```

```
## AIC BIC logLik
```

```
## 768.7183 802.8004 -375.3591
```

```
##
```

```
## Random effects:
```

```
## Formula: ~I(TopDress_kgha) | site_year
```

```
## Structure: General positive-definite, Log-Cholesky parametrization
```

```
## StdDev Corr
```

```
## (Intercept) 1.023988033 (Intr)
```

```
## I(TopDress_kgha) 0.003917507 -0.938
```

```
## Residual 0.711974429
```

```
##
## Fixed effects: GrainYield_Mgha ~ SI * TopDress_kgha + I(SI * SI)
##               Value Std.Error DF   t-value p-value
## (Intercept)   -11.61691  1.684234 321  -6.897447    0
## SI             39.64963  4.040201 321   9.813778    0
## TopDress_kgha   0.10523  0.016450 321   6.397171    0
## I(SI * SI)     -17.29309  2.477001 321  -6.981462    0
## SI:TopDress_kgha -0.10278  0.018162 321  -5.659094    0
## Correlation:
##               (Intr) SI      TpDrs_ I(SI*S
## SI              -0.957
## TopDress_kgha   -0.140  0.012
## I(SI * SI)       0.933 -0.993  0.074
## SI:TopDress_kgha 0.116 -0.013 -0.987 -0.075
##
## Standardized Within-Group Residuals:
##               Min           Q1           Med           Q3           Max
## -2.661180960 -0.649624675 -0.002562116  0.613738418  2.604603880
##
## Number of Observations: 331
## Number of Groups: 6
```

```
summary(sUAS_ndre_model_OR)$tTable
```

```
##               Value Std.Error DF   t-value   p-value
## (Intercept)   -11.6169128  1.68423378 321  -6.897447  2.824738e-11
## SI             39.6496313  4.04020072 321   9.813778  4.707103e-20
## TopDress_kgha   0.1052309  0.01644959 321   6.397171  5.593719e-10
## I(SI * SI)     -17.2930895  2.47700116 321  -6.981462  1.684349e-11
## SI:TopDress_kgha -0.1027811  0.01816211 321  -5.659094  3.374042e-08
```

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

```
## Analysis of Deviance Table (Type III tests)
```

```
##
```

```
## Response: GrainYield_Mgha
```

```
##               Chisq Df Pr(>Chisq)
## (Intercept)    47.575  1  5.295e-12 ***
## SI              96.310  1  < 2.2e-16 ***
## TopDress_kgha   40.924  1  1.583e-10 ***
## I(SI * SI)      48.741  1  2.921e-12 ***
## SI:TopDress_kgha 32.025  1  1.522e-08 ***
```

```
## ---
```

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

```
ndre_r_sq <- r.squaredGLMM(sUAS_ndre_model_OR)
```

```
ndre_r_sq_fixed <- round(ndre_r_sq[1] , digits = 2)
```

```
ndre_r_sq_fixed
```

```
## [1] 0.51
```

```
ndre_r_sq_total <- round(ndre_r_sq[2] , digits = 2)
```

```
ndre_r_sq_total
```

```
## [1] 0.82
```

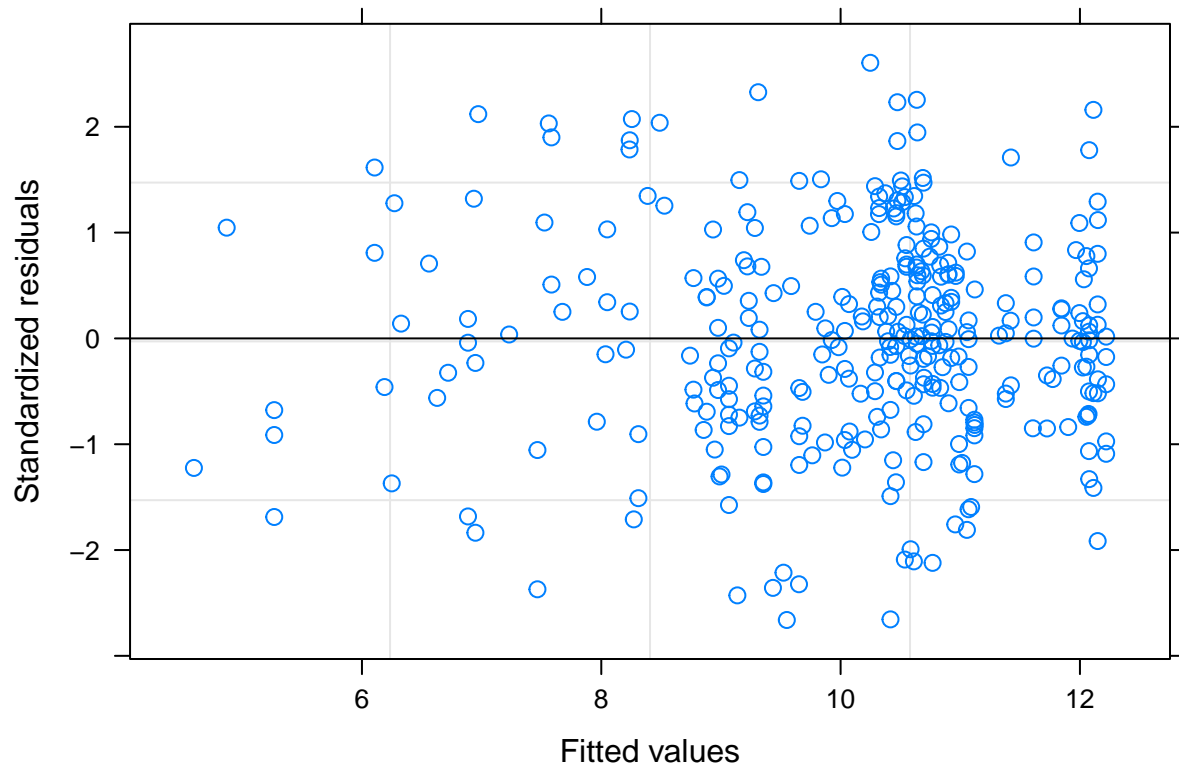


```
ndre_r_sq_random <- ndre_r_sq_total - ndre_r_sq_fixed  
ndre_r_sq_random
```

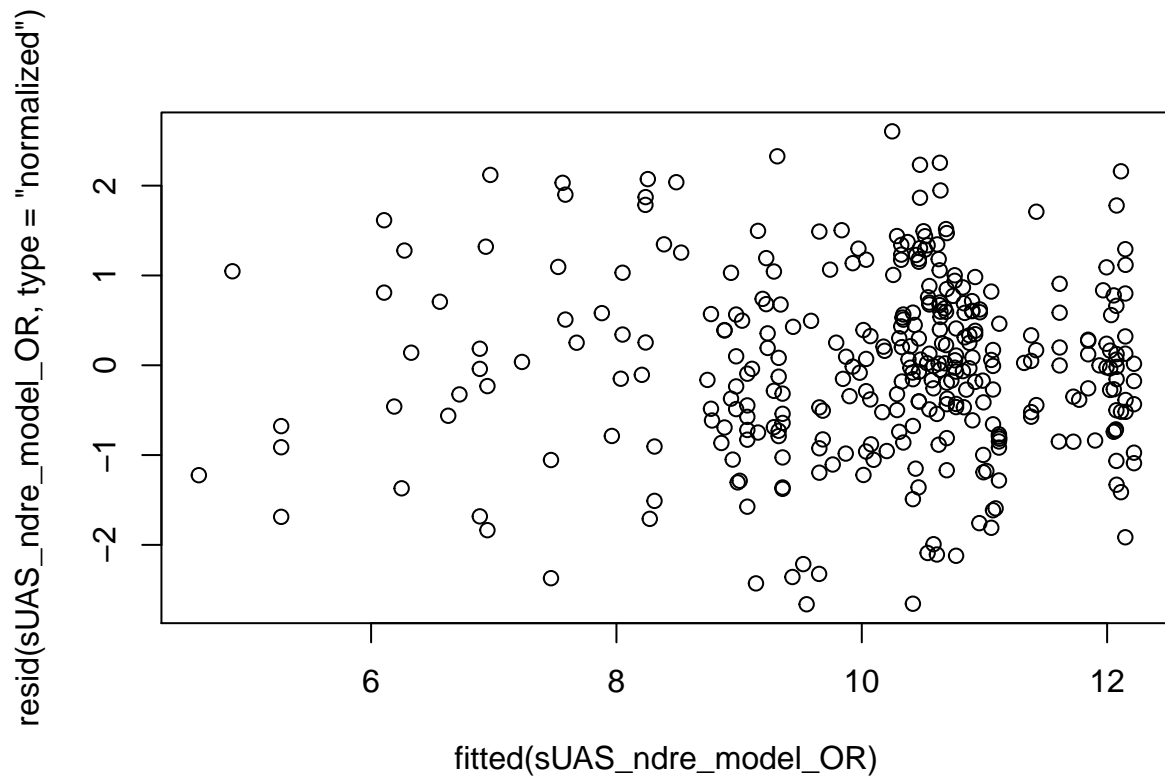
```
## [1] 0.31
```

model diagnostics

```
plot (sUAS_ndre_model_OR)
```



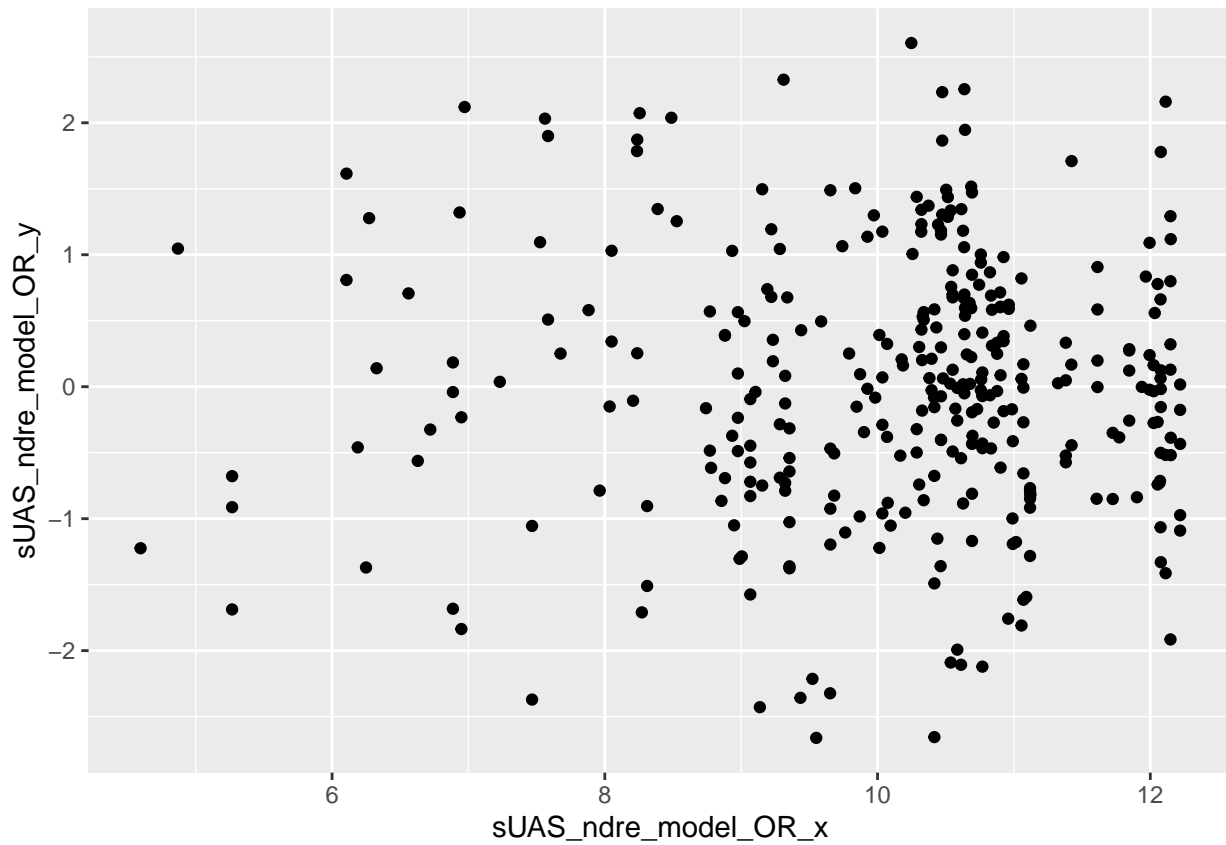
```
plot(resid(sUAS_ndre_model_OR, type = "normalized") ~fitted(sUAS_ndre_model_OR))
```



```
sUAS_ndre_model_OR_y <- resid(sUAS_ndre_model_OR, type = "normalized")
sUAS_ndre_model_OR_x <- fitted(sUAS_ndre_model_OR)

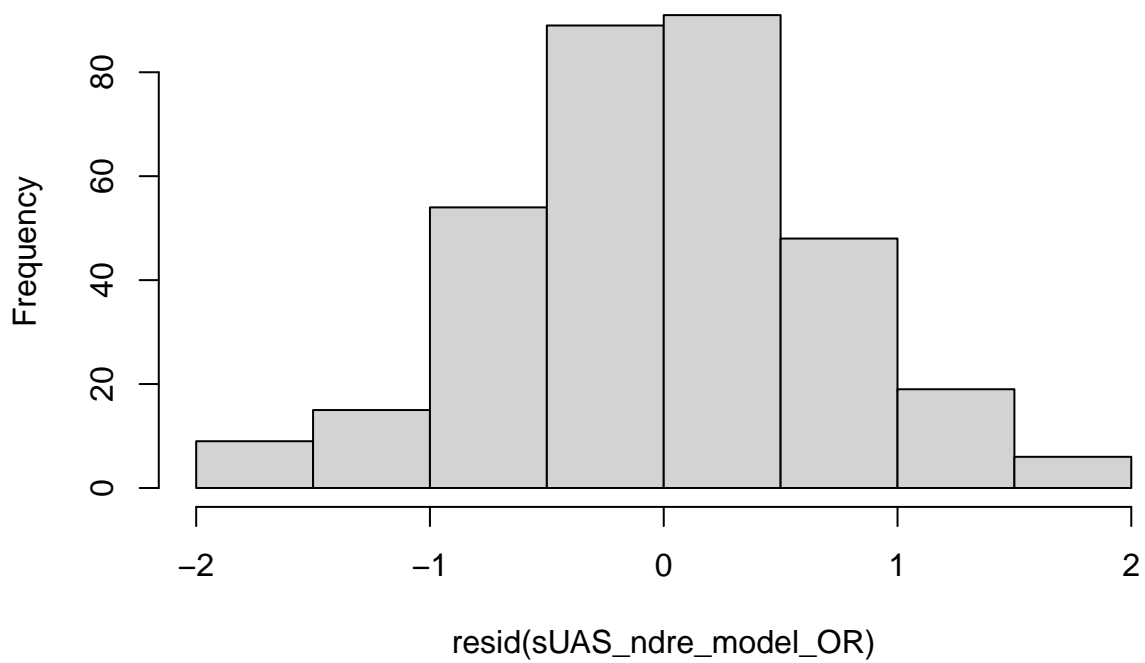
sUAS_ndre_model_ORresid_data <- data.frame(sUAS_ndre_model_OR_x , sUAS_ndre_model_OR_y)

ggplot( data = sUAS_ndre_model_ORresid_data , aes( x = sUAS_ndre_model_OR_x , y = sUAS_ndre_model_OR_y))
  geom_point(mapping = aes(sUAS_ndre_model_OR_x , sUAS_ndre_model_OR_y) , data = sUAS_ndre_model_ORresid_data)
```

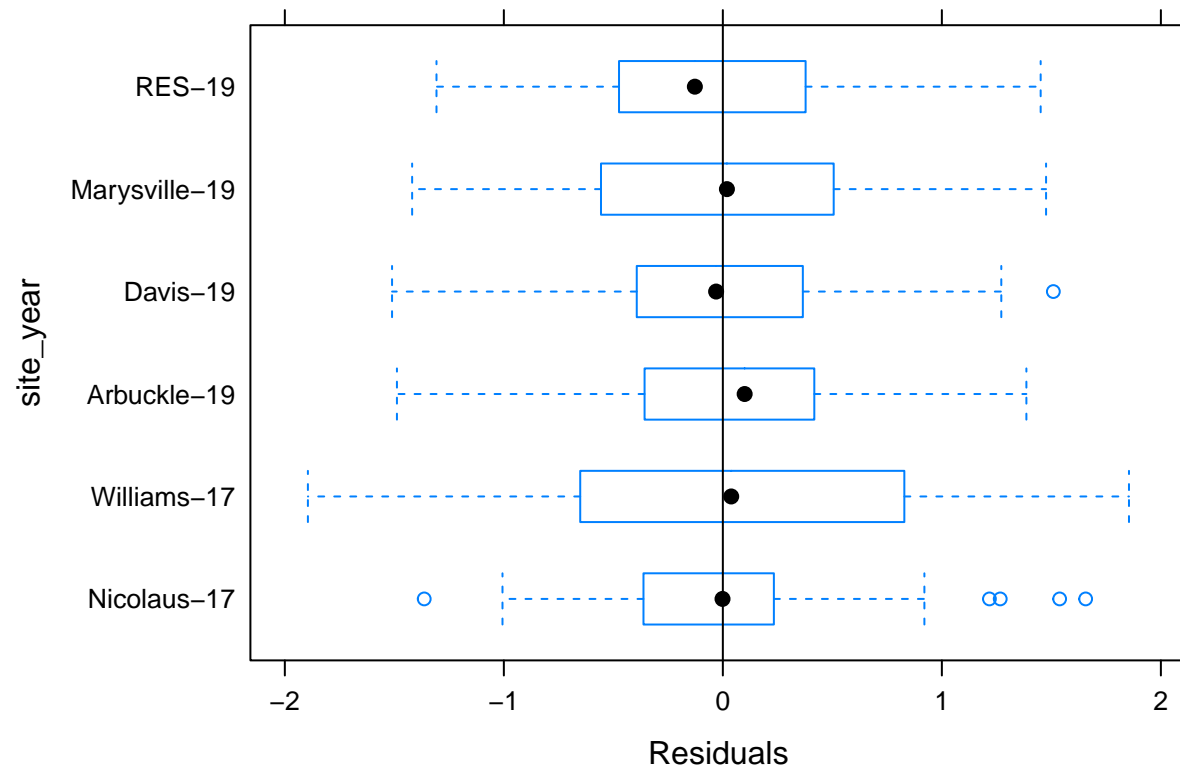


```
hist(resid(sUAS_ndre_model_OR))
```

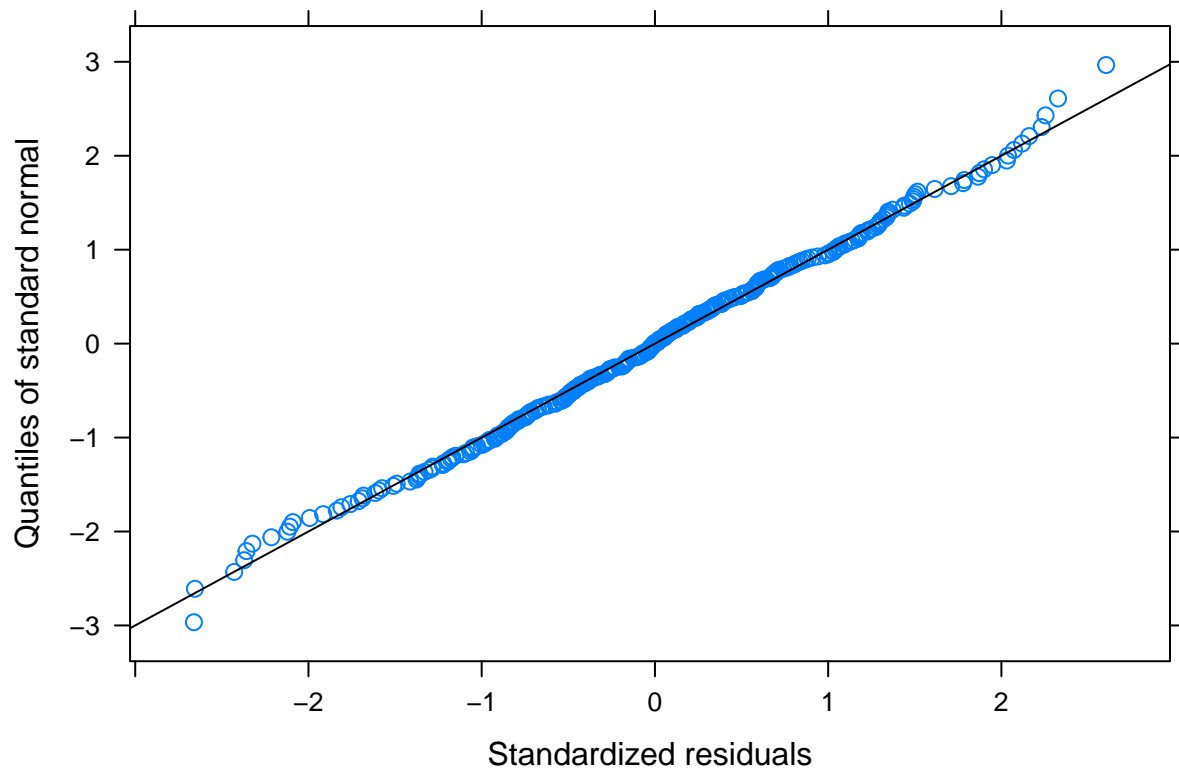
Histogram of resid(sUAS_ndre_model_OR)



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

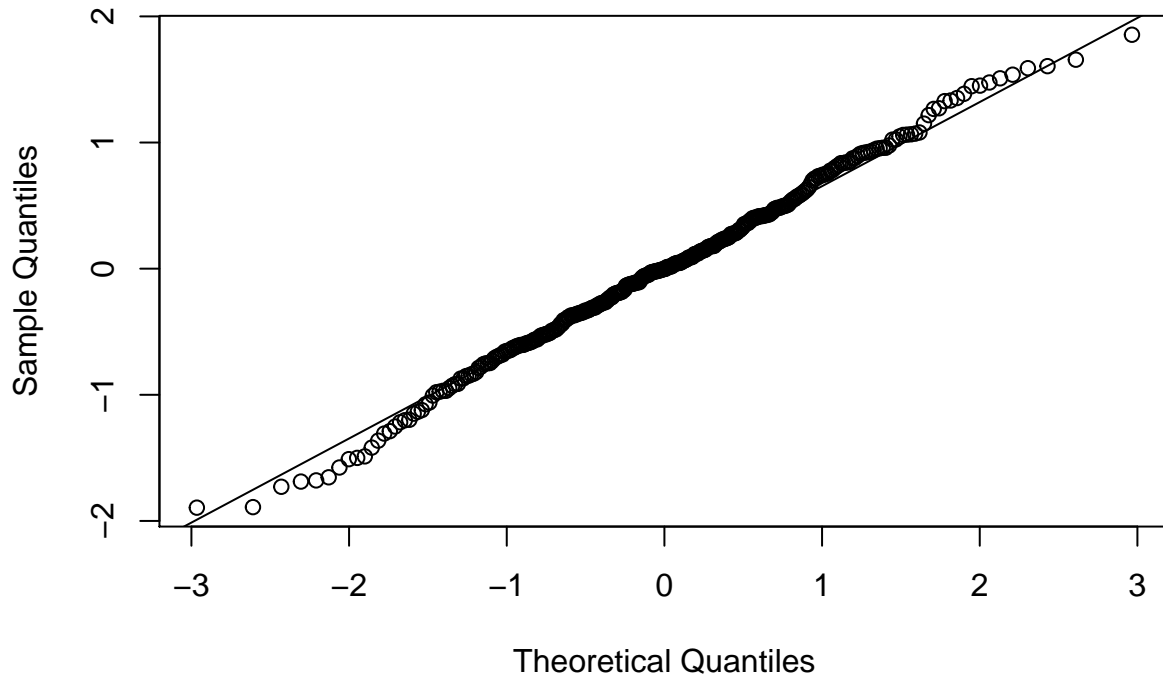


```
qqnorm(sUAS_ndre_model_OR, abline = c(0,1) )
```

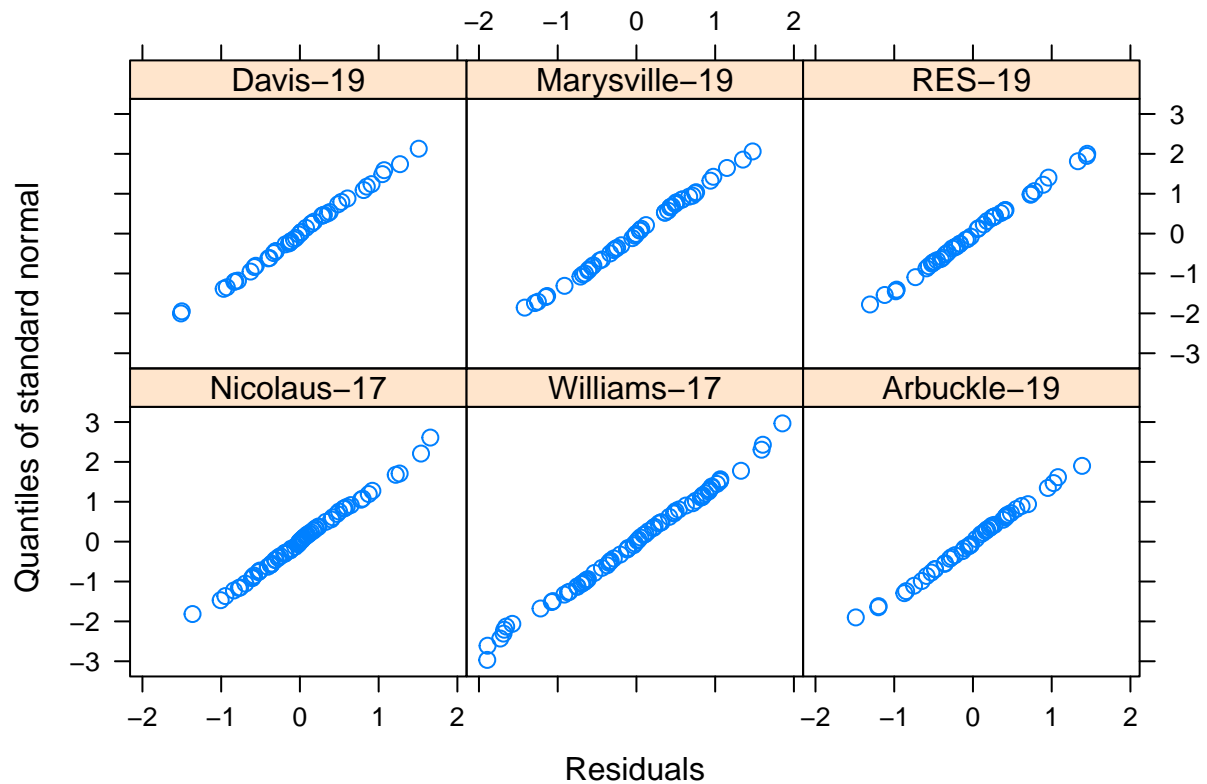


```
qqnorm(resid(sUAS_ndre_model_OR))
qqline(resid(sUAS_ndre_model_OR))
```

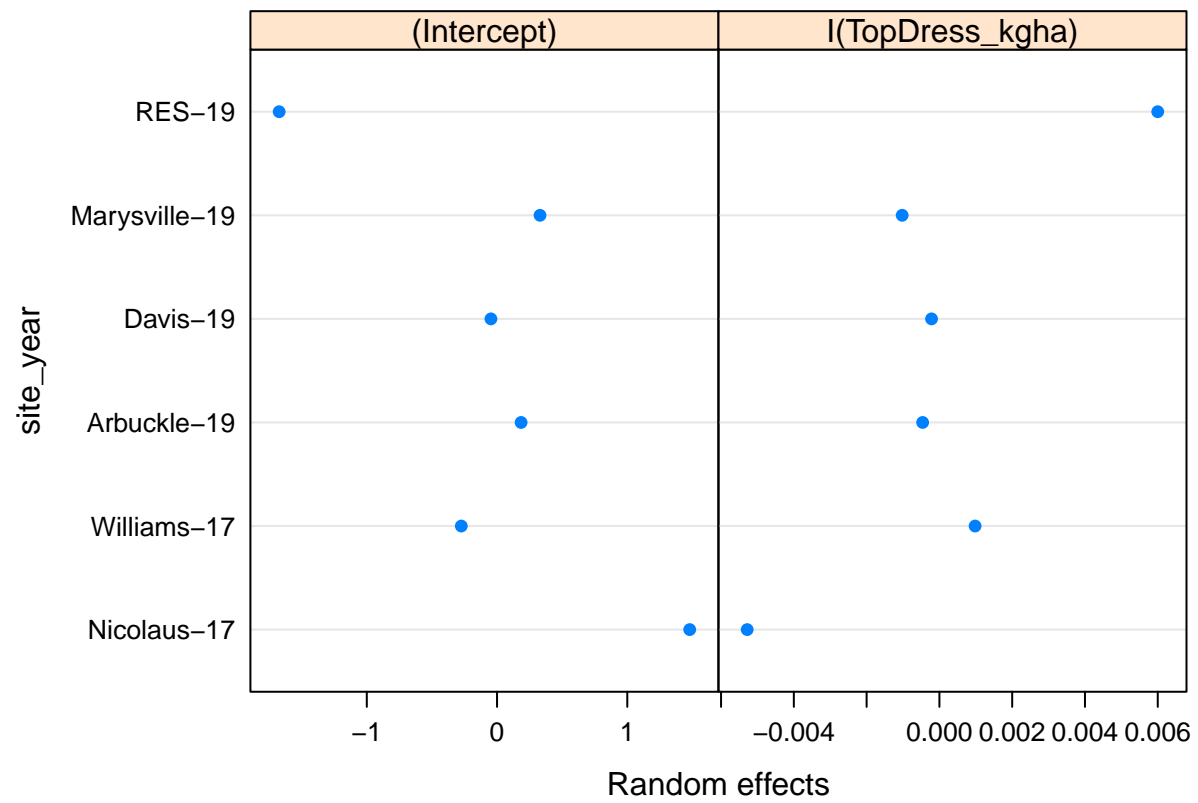
Normal Q-Q Plot



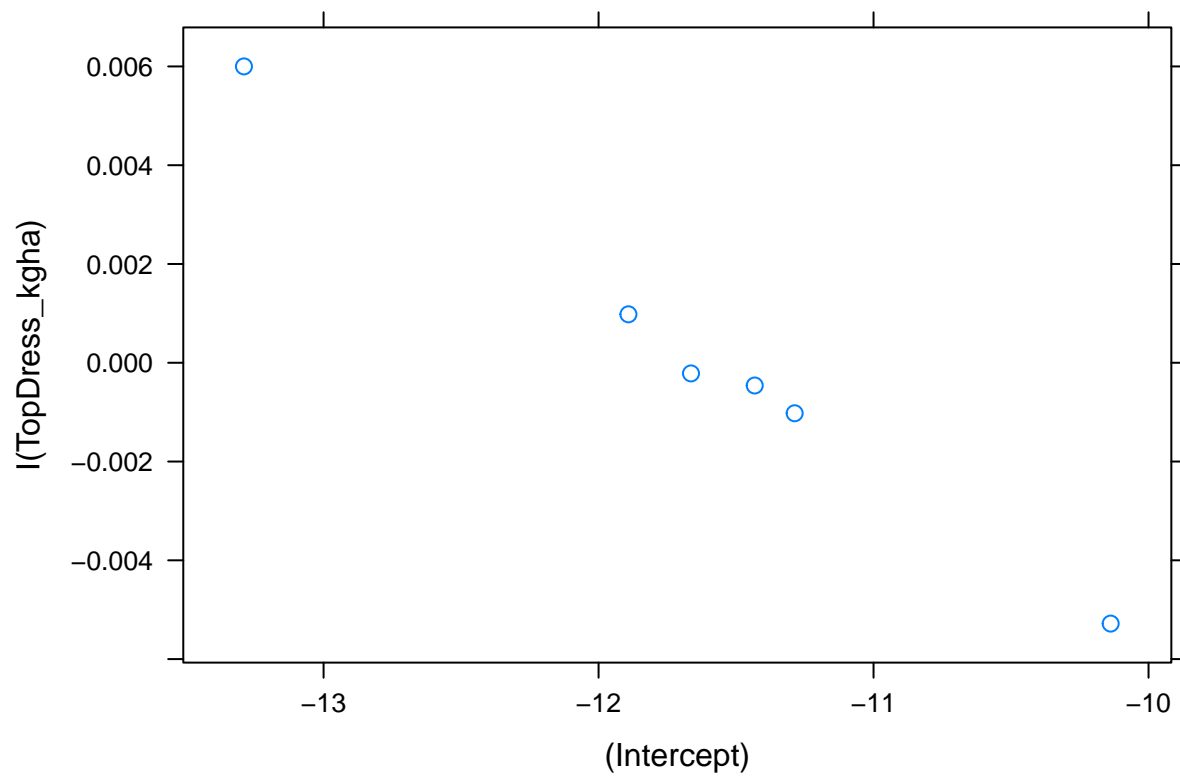
```
qqnorm(sUAS_ndre_model_OR , ~resid(.) | site_year)
```



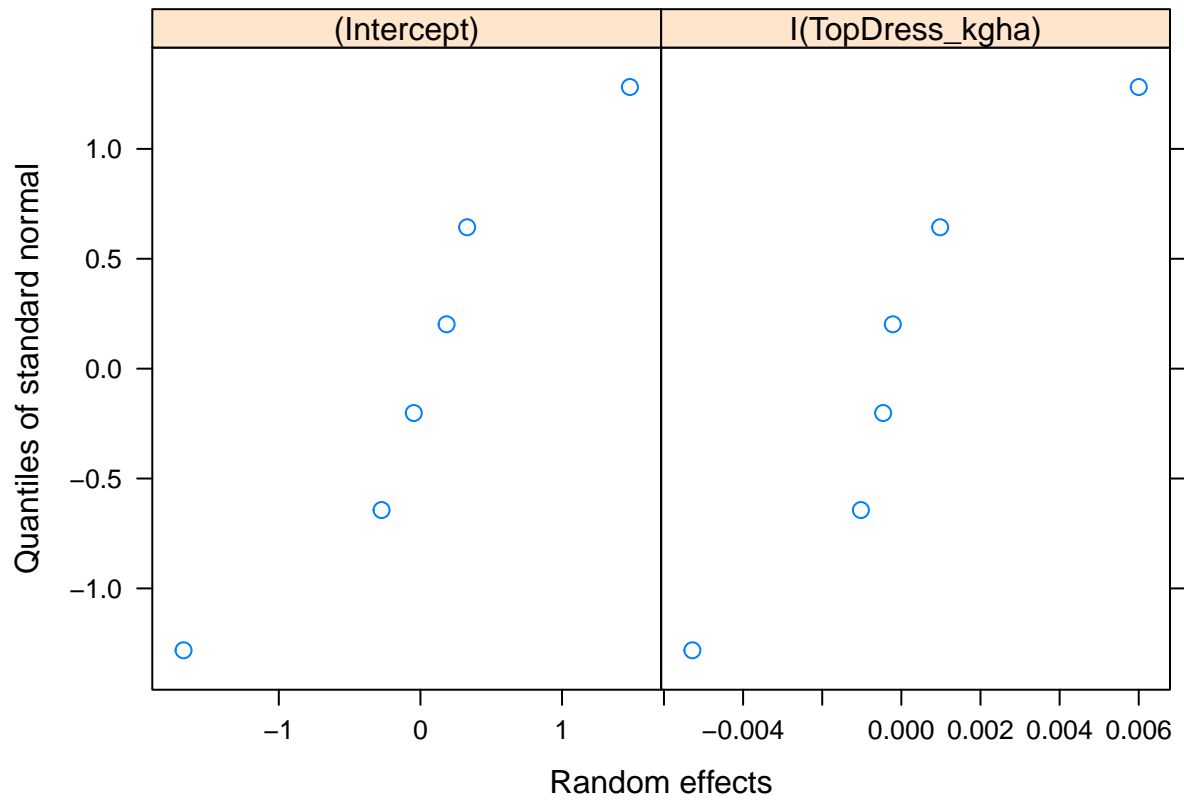
```
plot(ranef(sUAS_ndre_model_OR))
```



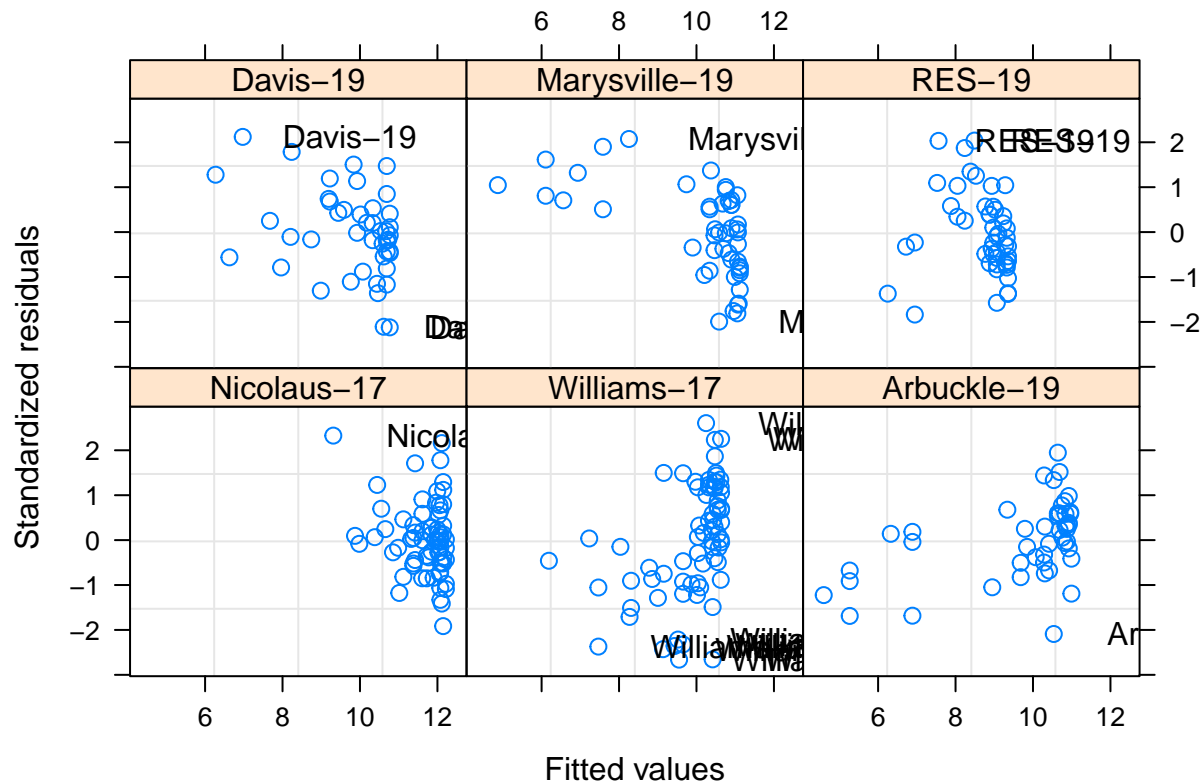
```
pairs(sUAS_ndre_model_OR , id = 0.1)
```



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



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



```
### emmeans
```

```
mylist <- list(SI=seq(0.528 , 1 , by = .0001), TopDress_kgha = c( 34 , 0))
```

```
sUAS_ndre_emmeans <- emmeans(sUAS_ndre_model_OR , ~TopDress_kgha * SI , at = mylist )
```

```
sUAS_ndre_emmeans_contrast <- as.data.frame(summary(contrast(sUAS_ndre_emmeans , "pairwise" , side = ">"))
```

```
sUAS_ndre_emmeans_contrast <- sUAS_ndre_emmeans_contrast %>%
  mutate(sUAS_ndre_Response_Index = 1 / SI )
```

```
sUAS_ndre_emmeans_contrast <- sUAS_ndre_emmeans_contrast %>%
  mutate(prob_postive_resp = (1 - p.value) * 100)
```

```
sUAS_ndre_emmeans_contrast <- sUAS_ndre_emmeans_contrast %>%
  mutate(t_score = abs((estimate - 0.26)) / SE)
```

```
sUAS_ndre_emmeans_contrast <- sUAS_ndre_emmeans_contrast %>%
  mutate(prob_greater_than_26 = if_else(estimate < 0.26 , pt(q = t_score , df = df , lower.tail = F) , 1))
```

```
sUAS_ndre_emmeans_contrast$sUAS_ndre_Response_Index_r <- (round(sUAS_ndre_emmeans_contrast$sUAS_ndre_Response_Index , 2))
```

```
sUAS_ndre_run1 <- 125 - 100
```

```
sUAS_ndre_rise1 <- 0.7890828 - 0.08455915
```

```
sUAS_ndre_slope1 <- round((sUAS_ndre_rise1 / sUAS_ndre_run1) * 5 , digits = 2)
```

```
sUAS_ndre_slope1
```

```
## [1] 0.14
```



```

sUAS_ndre_run2 <- 189 - 125
sUAS_ndre_rise2 <- 1.747235 - 0.7890828
sUAS_ndre_slope2 <- round((sUAS_ndre_rise2 / sUAS_ndre_run2) * 5 , digits = 3)
sUAS_ndre_slope2

## [1] 0.075

sUAS_ndre_mean_se <- sUAS_ndre_emmeans_contrast %>%
  select(SE) %>%
  summarise(mean_se = mean(SE))

sUAS_ndre_mean_se$mean_se <- round(sUAS_ndre_mean_se$mean_se , digits = 2)

sUAS_ndre_mean_se$mean_se

## [1] 0.14

###confidence interval

#at every 0.10 interval, what is the 90% confidence interval of the yield response, and what RI value d

str(sUAS_ndre_emmeans_contrast , give.attr = F)

## 'data.frame': 4721 obs. of 12 variables:
## $ contrast : Factor w/ 1 level "34 - 0": 1 1 1 1 1 1 1 1 1 1 ...
## $ SI : num 0.528 0.528 0.528 0.528 0.528 ...
## $ estimate : num 1.73 1.73 1.73 1.73 1.73 ...
## $ SE : num 0.243 0.243 0.243 0.243 0.243 ...
## $ df : num 321 321 321 321 321 321 321 321 321 321 ...
## $ t.ratio : num 7.12 7.12 7.12 7.12 7.13 ...
## $ p.value : num 3.47e-12 3.46e-12 3.46e-12 3.45e-12 3.45e-12 ...
## $ sUAS_ndre_Response_Index : num 1.89 1.89 1.89 1.89 1.89 ...
## $ prob_postive_resp : num 100 100 100 100 100 ...
## $ t_score : num 6.06 6.06 6.06 6.06 6.06 ...
## $ prob_greater_than_26 : num 1 1 1 1 1 ...
## $ sUAS_ndre_Response_Index_r: num 1.89 1.89 1.89 1.89 1.89 ...

sUAS_ndre_emmeans_contrast <- sUAS_ndre_emmeans_contrast %>%
  mutate(response_mgha_r = round(estimate , digits = 2),
         se_mgha_r = round(SE , digits = 3))

#estimate = 0.10 , SE = 0.112

print(round(0.10 - z_score*0.112 , digits = 2))

## [1] -0.08

print(round(0.10 + z_score*0.112 , digits = 2))

## [1] 0.28

#estimate = 0.20 , SE = 0.103

print(round(0.20 - z_score*0.103 , digits = 2))

## [1] 0.03

```

```

print(round(0.20 + z_score*0.103 , digits = 2))

## [1] 0.37
#estimate = 0.30 , SE = 0.097

print(round(0.30 - z_score*0.097 , digits = 2))

## [1] 0.14
print(round(0.30 + z_score*0.097 , digits = 2))

## [1] 0.46
#estimate = 0.40 , SE = 0.093

print(round(0.40 - z_score*0.093 , digits = 2))

## [1] 0.25
print(round(0.40 + z_score*0.093 , digits = 2))

## [1] 0.55
#estimate = 0.50 , SE = 0.093

print(round(0.50 - z_score*0.093 , digits = 2))

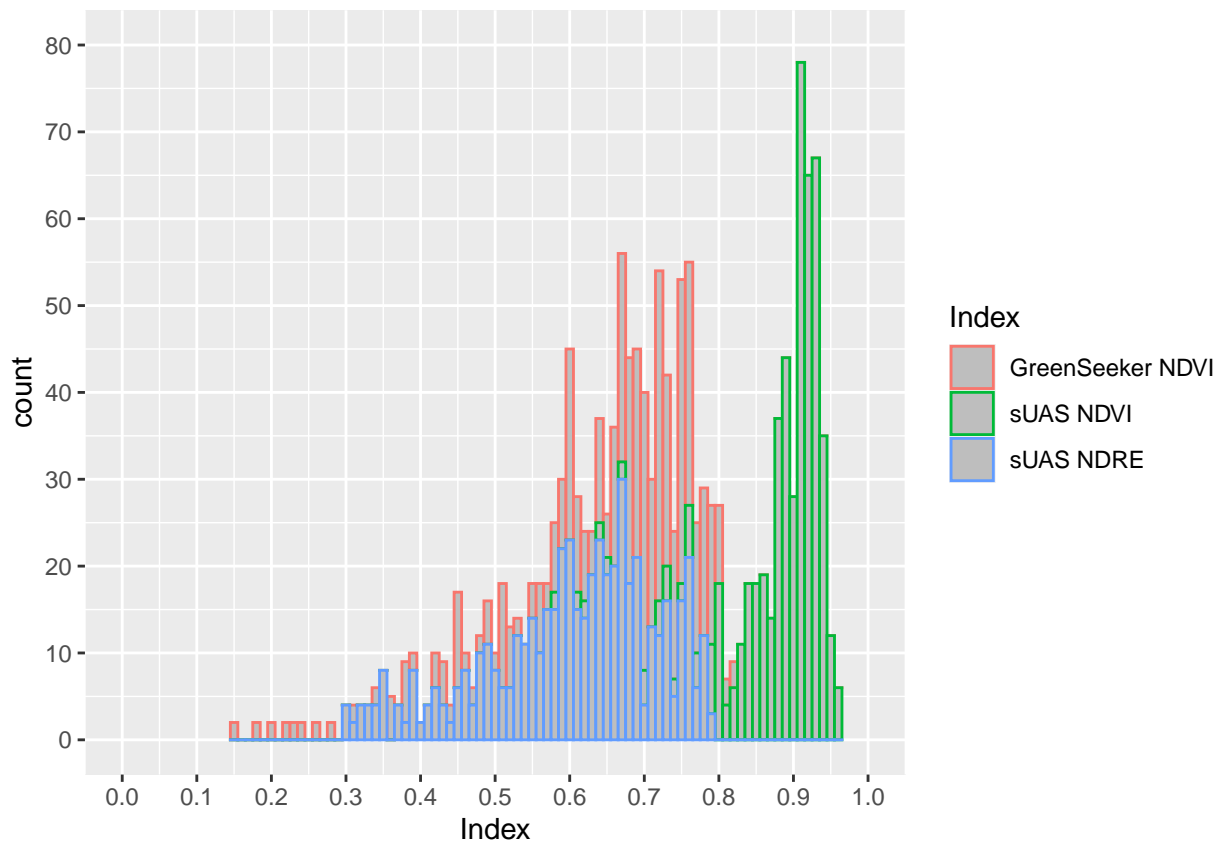
## [1] 0.35
print(round(0.50 + z_score*0.093 , digits = 2))

## [1] 0.65
#FIGURES
##FIGURE 2
hist_data <- paper3_data %>%
  select(Platform , Index) %>%
  group_by(Platform) %>%
  summarise(mean = mean(Index) , sd = sd(Index))

hist_data$mean = round(hist_data$mean , digits = 2)
hist_data$sd = round(hist_data$sd , digits = 2)

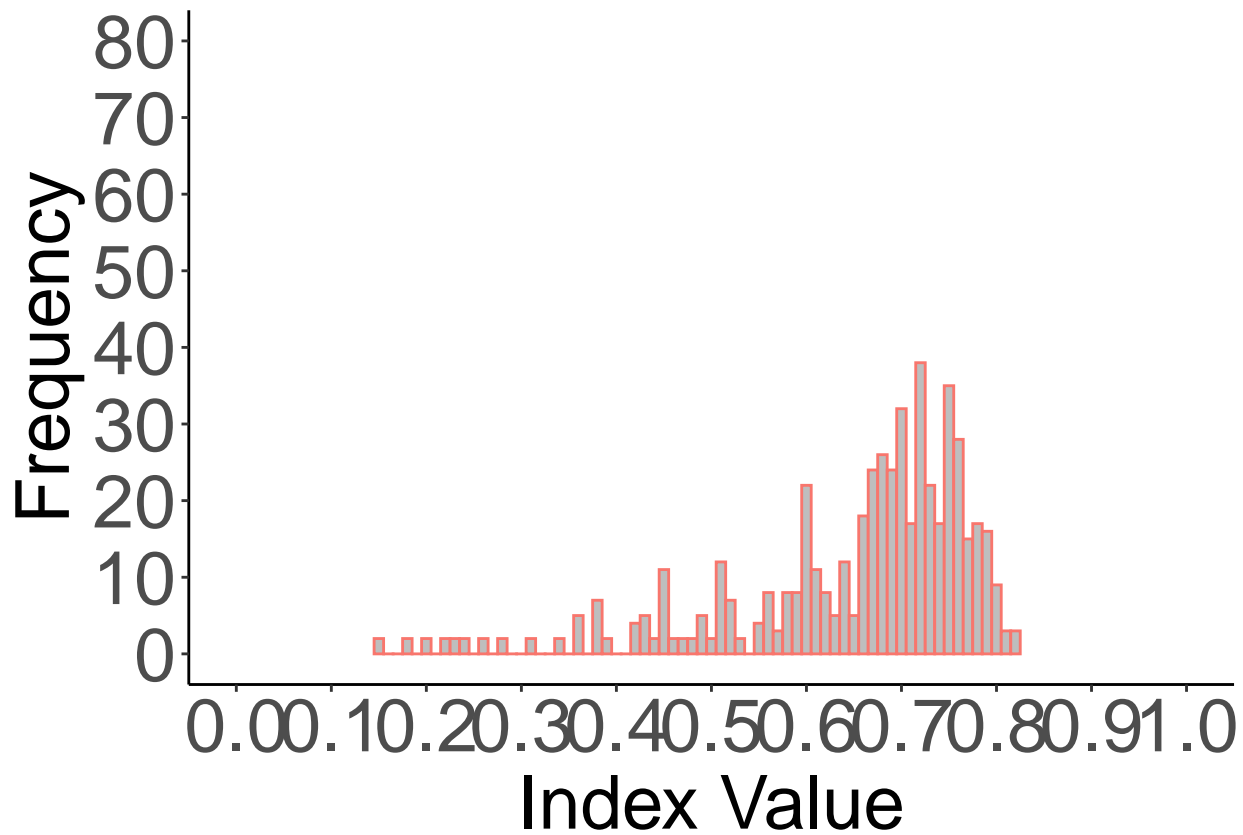
#This plot is for the legend
ggplot( data = paper3_data , aes(x = Index)) +
  geom_histogram( data = paper3_data , aes(x = Index , color = Platform) , binwidth = .01 , fill = "grey") +
  coord_cartesian(xlim = c(0 , 1) , ylim = c(0,80)) +
  scale_y_continuous(breaks = seq(0,100 , by = 10)) +
  scale_x_continuous(breaks = seq(0 , 1 , by = 0.1)) +
  scale_color_discrete( name = "Index" , labels = c("GreenSeeker NDVI" , "sUAS NDVI" , "sUAS NDRE"))

```



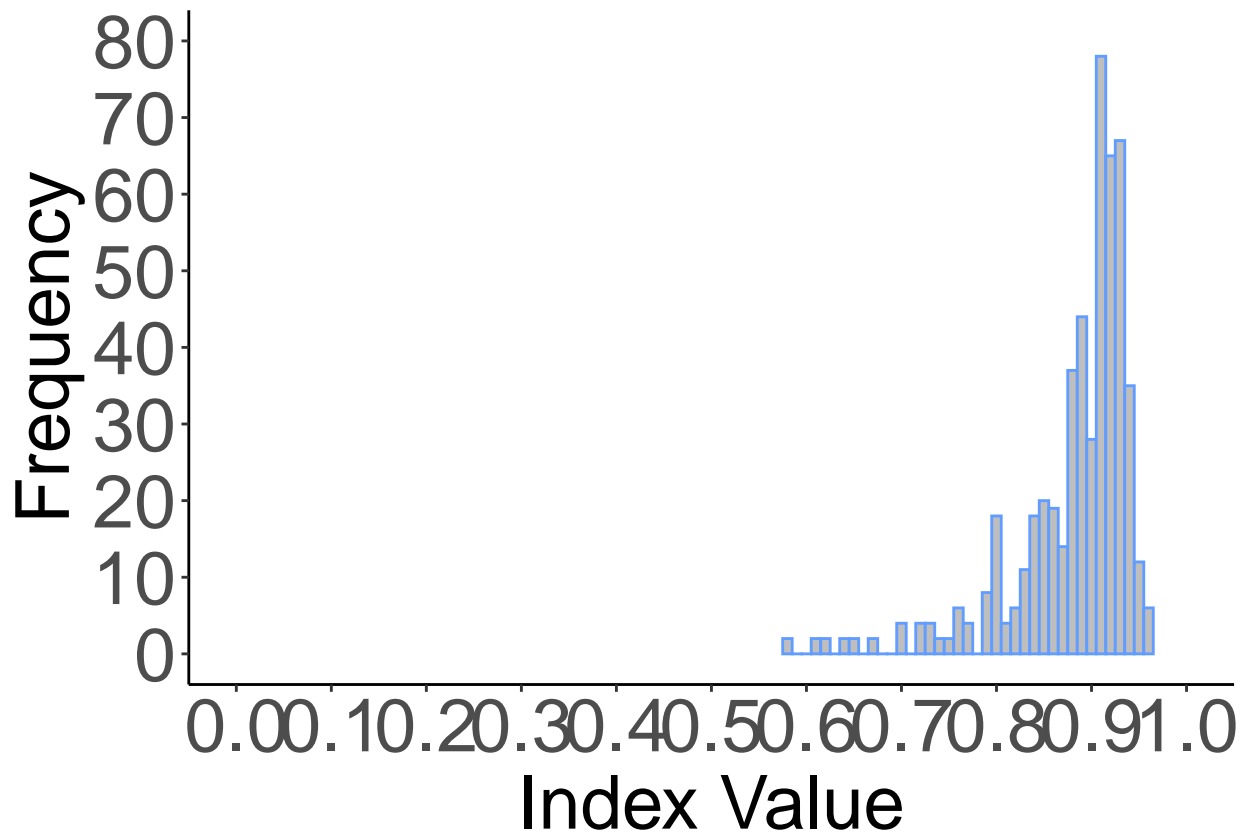
```
gs_ndvi_hist <- ggplot( data = paper3_gsdata , aes(x = Index)) +
  geom_histogram( data = paper3_gsdata , aes(x = Index , color = Platform) , binwidth = .01 , fill = "g
  theme_classic() +
  coord_cartesian(xlim = c(0 , 1) , ylim = c(0,80)) +
  scale_y_continuous(breaks = seq(0,100 , by = 10)) +
  scale_x_continuous(breaks = seq(0 , 1 , by = 0.1)) +
  theme(legend.position = "none") +
  labs(x = "Index Value" , y = "Frequency") +
  theme(axis.title = element_text(size = 28),
        axis.text = element_text(size = 28))

gs_ndvi_hist
```



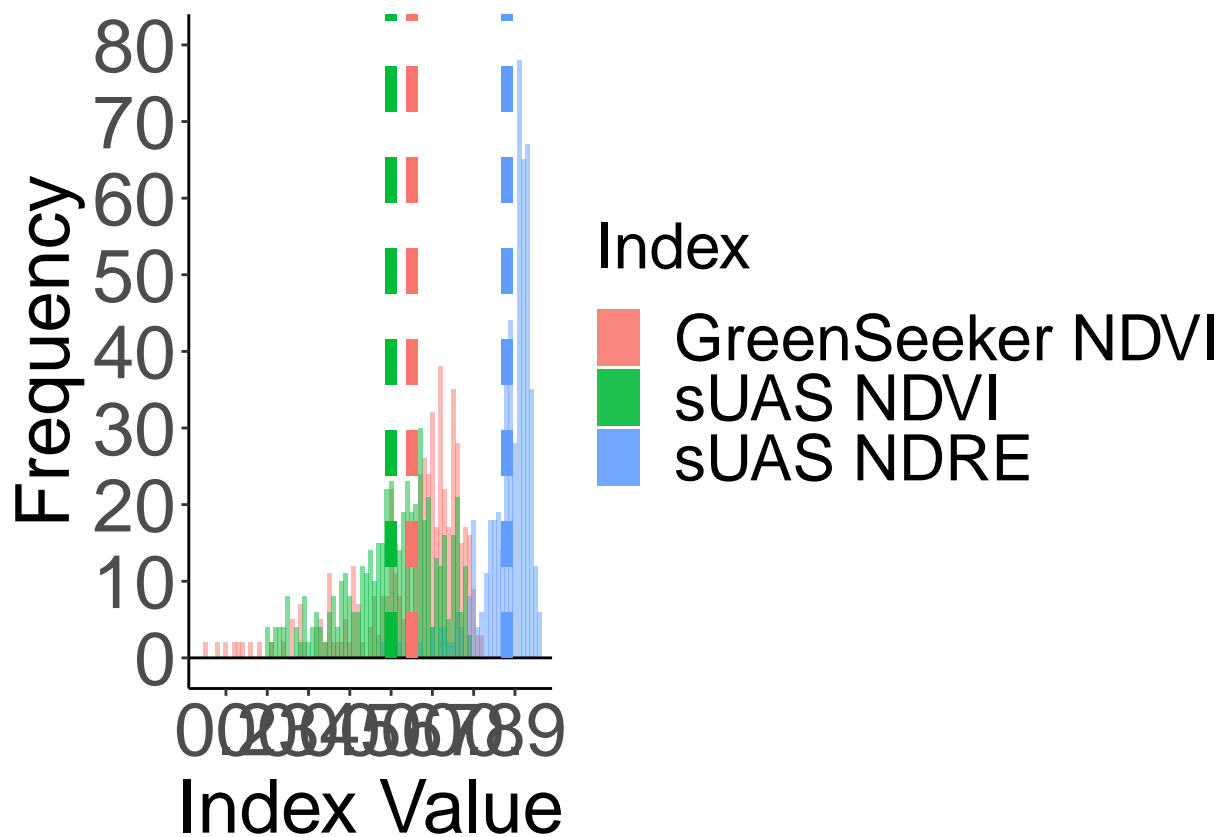
```
suas_ndvi_hist <- ggplot( data = paper3_uas_ndvi_data , aes(x = Index)) +
  geom_histogram( data = paper3_uas_ndvi_data , aes(x = Index ) , binwidth = .01 , fill = "grey" , color = "red" ) +
  theme_classic() +
  coord_cartesian(xlim = c(0 , 1) , ylim = c(0,80)) +
  scale_y_continuous(breaks = seq(0,100 , by = 10)) +
  scale_x_continuous(breaks = seq(0 , 1 , by = 0.1)) +
  theme(legend.position = "none") +
  labs(x = "Index Value" , y = "Frequency") +
  theme(axis.title = element_text(size = 28),
        axis.text = element_text(size = 28))

suas_ndvi_hist
```



```
Fig2 <- ggplot( data = paper3_data, aes( x = Index)) +
  geom_histogram(data = subset(paper3_data , Platform == 'GreenSeeker_NDVI') , aes(x = Index , fill = Platform)) +
  geom_histogram(data = subset(paper3_data , Platform == 'sUAS_NDVI') , aes(x = Index , fill = Platform)) +
  geom_histogram(data = subset(paper3_data , Platform == 'sUAS_NDRE') , aes(x = Index , fill = Platform)) +
  theme_classic() +
  coord_cartesian(xlim = c(0.15 , .95) , ylim = c(0,80)) +
  scale_y_continuous(breaks = seq(0,100 , by = 10)) +
  scale_x_continuous(breaks = seq(0 , 1 , by = 0.1)) +
  labs(x = "Index Value" , y = "Frequency") +
  theme(axis.title = element_text(size = 28),
        axis.text = element_text(size = 28),
        legend.text = element_text(size = 24),
        legend.title = element_text(size = 24)) +
  geom_hline(yintercept = 0) +
  scale_fill_discrete( name = "Index" , labels = c("GreenSeeker NDVI" , "sUAS NDVI" , "sUAS NDRE")) +
  geom_segment(aes(x = 0.65, y = 0, xend = 0.65, yend = Inf), linetype = "dashed" , size = 2 , alpha = 0.5) +
  geom_segment(aes(x = 0.88, y = 0, xend = 0.88, yend = Inf) ,linetype = "dashed" , size = 2 , alpha = 0.5) +
  geom_segment(aes(x = 0.60, y = 0, xend = 0.60, yend = Inf), linetype = "dashed" , size = 2 , alpha = 0.5)
```

Fig2



```
ggsave("FIGURES/Paper3_Fig_2.tiff" , Fig2 , compression = "lzw" , width = 16 , height = 12, type = "cairo")
```

```
##FIGURE 3
```

```
platform <- data.frame(
  label = c("NDVI[GS]" ,
            "NDVI[UAS]" ,
            "NDRE[UAS]"
  ) ,
  Platform = c("GreenSeeker_NDVI" ,
               "sUAS_NDVI" ,
               "sUAS_NDRE"
  ) ,
  x_position = c(Inf , Inf , Inf ) ,
  x_just_var = c( 2.5 , 2.2 , 2 ) ,
  y_position = c(-Inf , -Inf , -Inf ) ,
  y_just_var = c(-12 , -12 , -12)
)
```

```
equation <- data.frame(
  label = c(gs_ndvi_qm_eqn,
            sUAS_ndvi_qm_eqn,
            sUAS_ndre_qm_eqn
  ) ,
  Platform = c("GreenSeeker_NDVI" ,
               "sUAS_NDVI" ,
               "sUAS_NDRE"
  ) ,

```

```

x_position = c(Inf , Inf , Inf ),
x_just_var = c(1.05 , 1.05 , 1.05) ,
y_position = c(-Inf , -Inf , -Inf ),
y_just_var = c(-8 , -8 , -8)
)

rsquared <- data.frame(
  label = c("R^2 == '0.80'",
            "R^2 == '0.80'" ,
            "R^2 == '0.82'"
            ) ,
  Platform = c("GreenSeeker_NDVI" ,
               "sUAS_NDVI",
               "sUAS_NDRE"
               ),
  x_position = c(Inf , Inf , Inf ),
  x_just_var = c(2.25 , 2.25 , 2) ,
  y_position = c(-Inf , -Inf , -Inf ),
  y_just_var = c(-8.5 , -8.5 , -8.5)
)

asymptote <- data.frame(
  label = c("N[UP] == 173~kg~N~ha^-1",
            "N[UP] == 164~kg~N~ha^-1" ,
            "N[UP] == 208~kg~N~ha^-1"
            ) ,
  Platform = c("GreenSeeker_NDVI" ,
               "sUAS_NDVI",
               "sUAS_NDRE"
               ),
  x_position = c(Inf , Inf , Inf ),
  x_just_var = c(1.85 , 1.85 , 1.65) ,
  y_position = c(-Inf , -Inf , -Inf ),
  y_just_var = c(-7.75 , -7.75 , -7.75)
)

platform$Platform = factor(platform$Platform, levels=c("GreenSeeker_NDVI" , "sUAS_NDVI" , "sUAS_NDRE"))
equation$Platform = factor(equation$Platform, levels=c("GreenSeeker_NDVI" , "sUAS_NDVI" , "sUAS_NDRE"))
rsquared$Platform = factor(rsquared$Platform, levels=c("GreenSeeker_NDVI" , "sUAS_NDVI" , "sUAS_NDRE"))
asymptote$Platform = factor(asymptote$Platform, levels=c("GreenSeeker_NDVI" , "sUAS_NDVI" , "sUAS_NDRE"))
gs_ndvi_qm_df$Platform = as.factor(gs_ndvi_qm_df$Platform)
sUAS_ndvi_qm_df$Platform = as.factor(sUAS_ndvi_qm_df$Platform)
sUAS_ndre_qm_df$Platform = as.factor(sUAS_ndre_qm_df$Platform)
asym_data_gs_ndvi <- data.frame(x = gs_ndvi_qm_sym_x , y = gs_ndvi_qm_sym_y , xend = gs_ndvi_qm_sym_x ,
asym_data_sUAS_ndvi <- data.frame(yint = sUAS_ndvi_qm_sym_y , Platform = factor("sUAS_NDVI"))
asym_data_sUAS_ndre <- data.frame(yint = sUAS_ndre_qm_sym_y , Platform = factor("sUAS_NDRE"))

label_text <- data.frame(
  label = c("A" ,
            "B",
            "C"
            ) ,
  Platform = factor(c("GreenSeeker_NDVI" ,

```

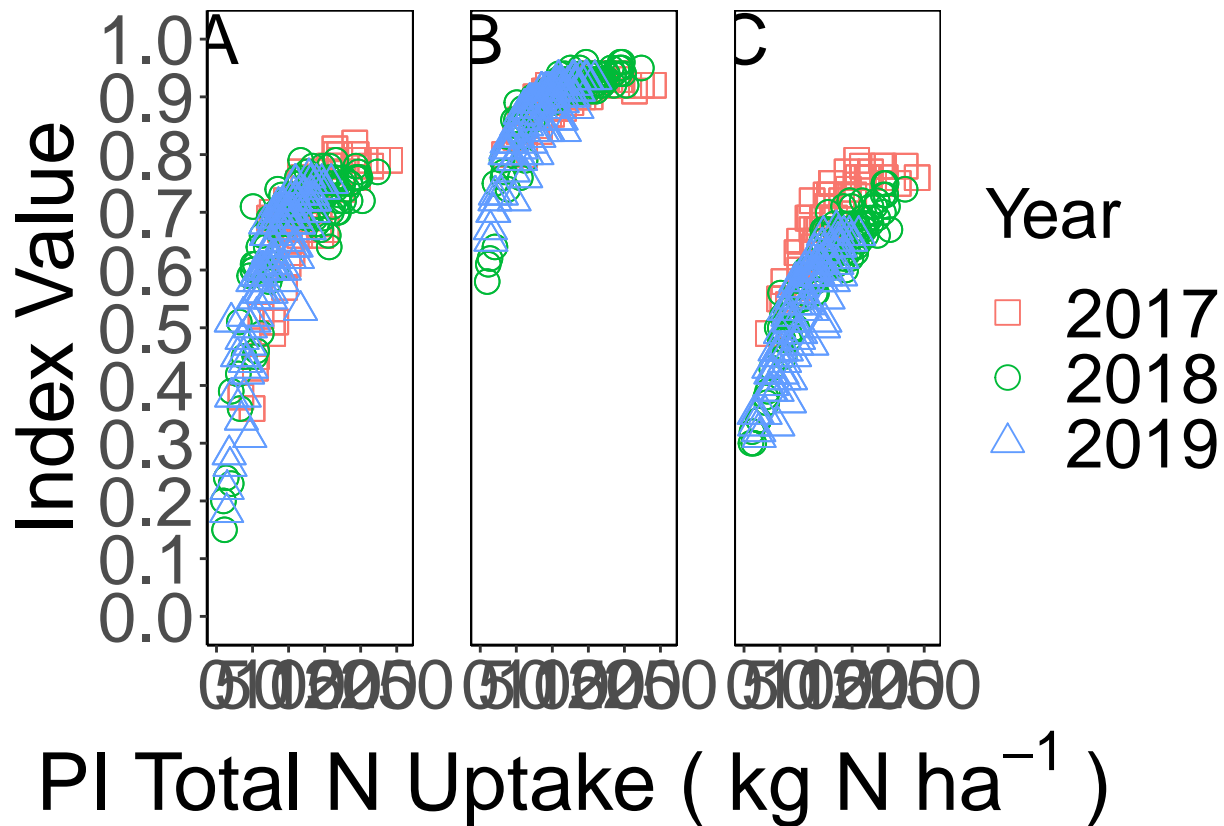
```

      "sUAS_NDVI" ,
      "sUAS_NDRE"
    )),
  x_position = c(0 , 0 , 0 ) ,
  y_position = c(1 , 1 , 1 ))

Nup_fig <- ggplot( data = paper3_data , aes ( x = PI_N_Uptake , y = Index )) +
  geom_point( mapping = aes( x = PI_N_Uptake ,
                             y = Index ,
                             shape = year,
                             color = year) ,
              size = 4 ,
              data = paper3_data) +
  labs( x = "PI Total N Uptake ( kg N ha-1~)" ,
        y = "Index Value",
        shape = "Year",
        color = "Year") +
  facet_grid(~Platform ) +
  theme_classic() +
  theme(axis.title = element_text(size = 30),
        axis.text = element_text(size = 26),
        legend.text = element_text(size = 26),
        legend.title = element_text(size = 26),
        #legend.position = "none",
        panel.spacing = unit(1.5, "lines"),
        panel.background = element_rect(fill = "white", color = "grey0"),
        strip.text.x = element_blank()) +
  coord_cartesian(ylim = c(0 , 1) , xlim = c(0, 260)) +
  scale_y_continuous(breaks = seq(0 , 1 , by = .1)) +
  scale_x_continuous(breaks = seq(0 , 250, by = 50)) +
  scale_shape_manual(values = c(0 , 1 , 2)) +
  geom_text( data = label_text ,
            mapping = aes( x = x_position ,
                           y = y_position ,
                           label = label) ,
            size = 9 ,
            parse = T)

```

Nup_fig



```
Fig3 <- Nup_fig +
  geom_line(data = gs_ndvi_qm_df , aes(x = gsndvi_df$PI_N_Uptake , y = gs_ndvi_fit_qm ) , size = 2 , col = "green") +
  geom_segment(data = asym_data_gs_ndvi , aes(x = x, y = y, xend = xend , yend = yend), linetype = "dashed", size = 2 , col = "red") +
  geom_line(data = sUAS_ndvi_qm_df , aes(x = sUASndvi_df$PI_N_Uptake , y = sUAS_ndvi_fit_qm ) , size = 2 , col = "blue") +
  geom_line(data = sUAS_ndre_qm_df , aes(x = sUASndre_df$PI_N_Uptake , y = sUAS_ndre_fit_qm ) , size = 2 , col = "blue") +
  geom_text( data = platform ,
    mapping = aes( x = x_position ,
      hjust = x_just_var ,
      y = y_position ,
      vjust = y_just_var ,
      label = label) ,
    size = 6 ,
    parse = T) +
  geom_text( data = equation ,
    mapping = aes( x = x_position ,
      hjust = x_just_var ,
      y = y_position ,
      vjust = y_just_var ,
      label = label) ,
    size = 6 ,
    parse = T) +
  geom_text( data = rsquared ,
    mapping = aes( x = x_position ,
      hjust = x_just_var ,
      y = y_position ,
      vjust = y_just_var ,
      label = label) ,
    size = 6 ,
    parse = T)
```

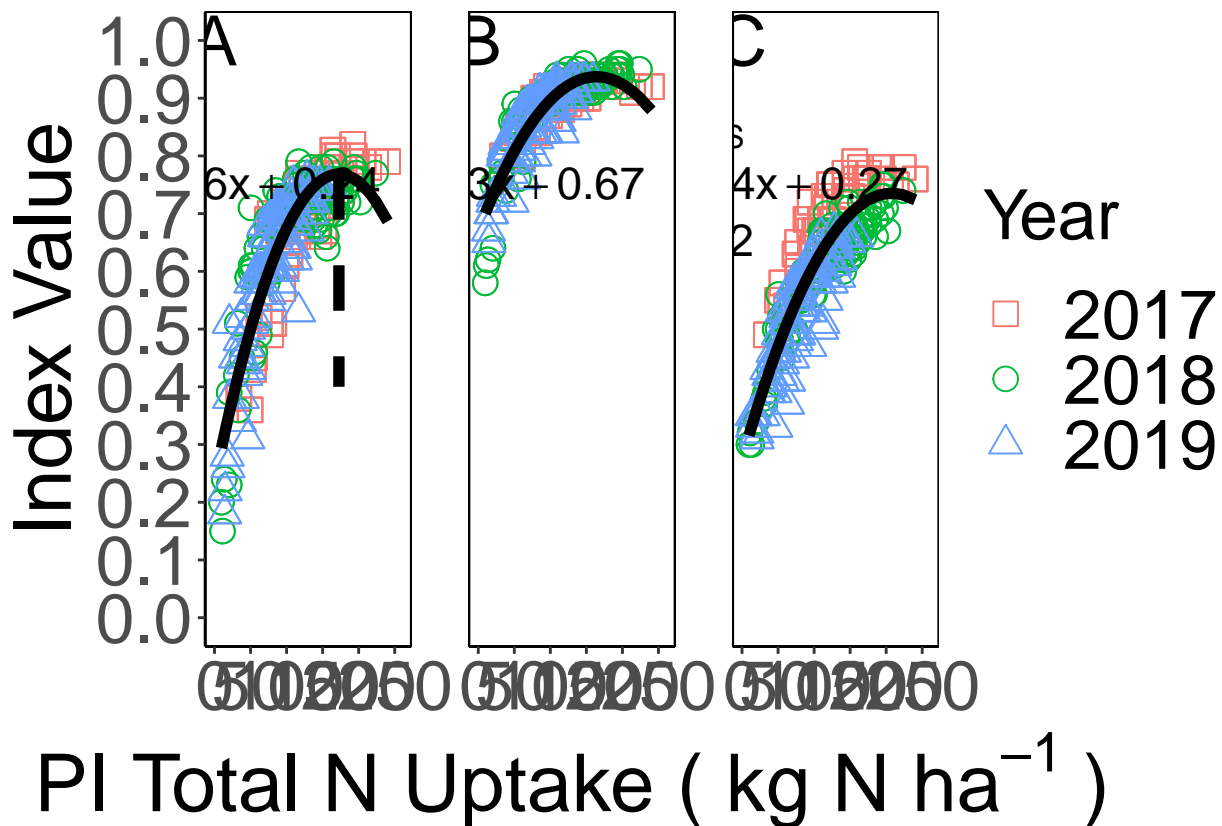
```

      parse = T) +
  geom_text( data = asymptote ,
            mapping = aes( x = x_position ,
                          hjust = x_just_var ,
                          y = y_position ,
                          vjust = y_just_var ,
                          label = label) ,

            size = 6 ,
            parse = T)

```

Fig3



```

ggsave("FIGURES/Paper3_Fig_3.tiff" , Fig3 , compression = "lzw" , width = 19 , height = 9.5, type = "cairo")

```

##FIGURE 4

```

estimated_yield_response <- seq(0.1 , 0.5 , by = 0.1)
index <- as.factor("GreenSeeker NDVI")
RI <- c('1.000' , '1.037' , '1.078' , '1.123' , '1.172')
lower_limit <- c('1.000' , '1.000' , '1.010' , '1.049' , '1.091')
upper_limit <- c('1.078' , '1.114' , '1.157' , '1.208' , '1.265')
position <- seq(0.08 , 0.48 , by = 0.1)

ci_plot_data1 <- data.frame(index ,
                           estimated_yield_response ,
                           RI ,
                           lower_limit ,
                           upper_limit ,

```

```

        position)

estimated_yield_response <- seq(0.1 , 0.5 , by = 0.1)
index <- as.factor("sUAS NDVI")
RI <- c(NA , '1.000' , '1.013' , '1.028' , '1.044')
lower_limit <- c(NA , '1.000' , '1.000' , '1.005' , '1.019' )
upper_limit <- c(NA , '1.027' , '1.039' , '1.053' , '1.069' )
position <- seq(0.1 , 0.5 , by = 0.1)

ci_plot_data2 <- data.frame(index ,
                           estimated_yield_response ,
                           RI ,
                           lower_limit ,
                           upper_limit ,
                           position)

estimated_yield_response <- seq(0.1 , 0.5 , by = 0.1)
index <- as.factor("sUAS NDRE")
RI <- c('1.003' , '1.032' , '1.064' , '1.097' , '1.132')
lower_limit <- c('1.000' , '1.000' , '1.015' , '1.048' , '1.080' )
upper_limit <- c('1.057' , '1.087' , '1.118' , '1.150' , '1.189' )
position <- seq(0.12 , 0.52 , by = 0.1)

ci_plot_data3 <- data.frame(index ,
                           estimated_yield_response ,
                           RI ,
                           lower_limit ,
                           upper_limit ,
                           position)

ci_plot_data <- rbind(ci_plot_data1 , ci_plot_data2 , ci_plot_data3)

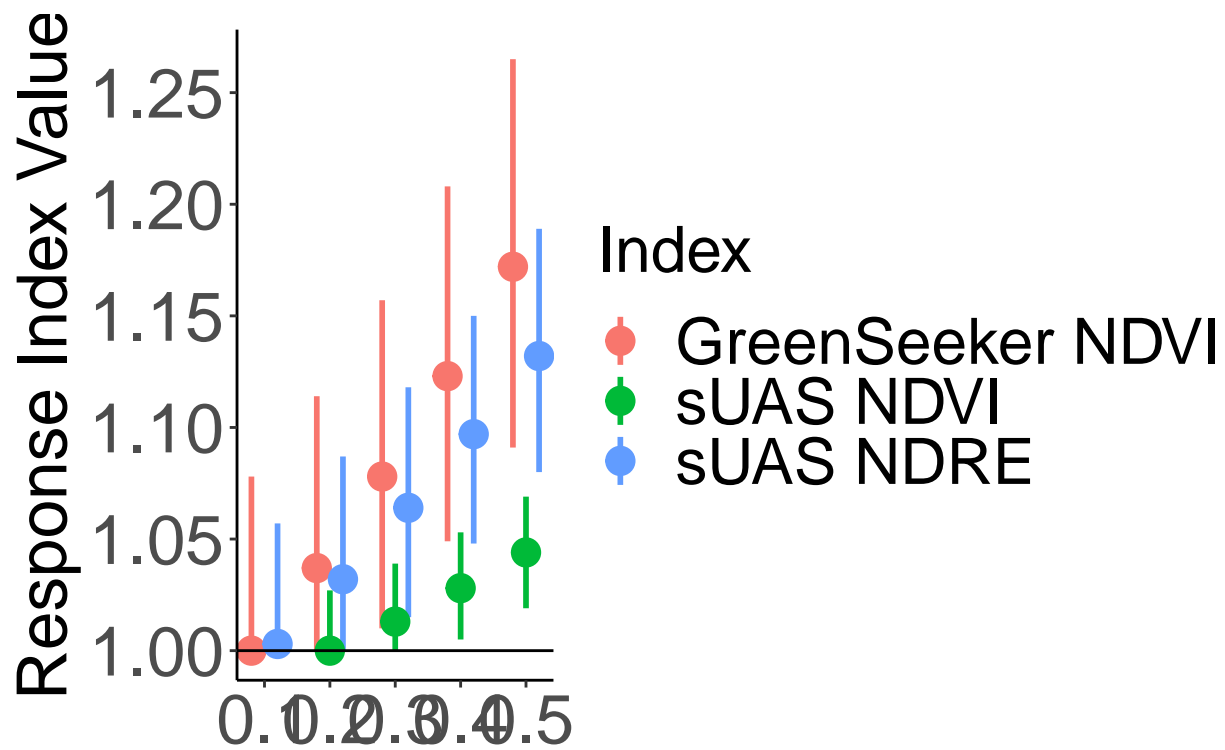
ci_plot_data$RI <- as.numeric(ci_plot_data$RI)
ci_plot_data$lower_limit <- as.numeric(ci_plot_data$lower_limit)
ci_plot_data$upper_limit <- as.numeric(ci_plot_data$upper_limit)

Fig4 <- ggplot(data = ci_plot_data , aes (y = RI , x = estimated_yield_response)) +
  geom_pointrange( data = ci_plot_data , aes(y = RI , x = position , ymin = lower_limit , ymax = upper_limit)) +
  theme_classic() +
  scale_y_continuous(breaks = seq(1 , 1.28 , by = 0.05)) +
  geom_hline(yintercept = 1) +
  labs(x = "Estimated Grain Yield Response ( Mg ha-1~") , y = "Response Index Value" , color = "Index") +
  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)
  )

```

Fig4

```
## Warning: Removed 1 rows containing missing values (geom_pointrange).
```



d Grain Yield Response (Mg ha⁻¹)

```
ggsave("FIGURES/Paper3_Fig_4.tiff" , Fig4 , compression = "lzw" , width = 15 , height = 12, type = "cairo")
```

```
## Warning: Removed 1 rows containing missing values (geom_pointrange).
```