

Dengue.R

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```
#Dengue Analysis
#Created by Ratnam Dubey
#https://www.drivendata.org/competitions/44/dengai-predicting-disease-
spread/submissions/
# Load Libraries

pkgs <- c('tidyverse','scales' , 'corrplot', 'magrittr','corrplot' , 'zoo',
'RColorBrewer', 'gridExtra', 'MASS', 'plyr' , 'dplyr' , 'plotly' )
invisible(lapply(pkgs, require, character.only = T))

## Loading required package: tidyverse

## Warning in library(package, lib.loc = lib.loc, character.only = TRUE,
## logical.return = TRUE, : there is no package called 'tidyverse'

## Loading required package: scales

## Warning: package 'scales' was built under R version 3.3.3

## Loading required package: corrplot

## Warning: package 'corrplot' was built under R version 3.3.3

## Loading required package: magrittr

## Loading required package: zoo

## Warning: package 'zoo' was built under R version 3.3.3

##
## Attaching package: 'zoo'

## The following objects are masked from 'package:base':
##
##   as.Date, as.Date.numeric

## Loading required package: RColorBrewer

## Loading required package: gridExtra

## Loading required package: MASS

## Loading required package: plyr

## Loading required package: dplyr
```

```
##
## Attaching package: 'dplyr'

## The following objects are masked from 'package:plyr':
##
##   arrange, count, desc, failwith, id, mutate, rename, summarise,
##   summarize

## The following object is masked from 'package:MASS':
##
##   select

## The following object is masked from 'package:gridExtra':
##
##   combine

## The following objects are masked from 'package:stats':
##
##   filter, lag

## The following objects are masked from 'package:base':
##
##   intersect, setdiff, setequal, union

## Loading required package: plotly
## Warning: package 'plotly' was built under R version 3.3.3
## Loading required package: ggplot2
## Warning: package 'ggplot2' was built under R version 3.3.3

##
## Attaching package: 'plotly'

## The following object is masked from 'package:ggplot2':
##
##   last_plot

## The following objects are masked from 'package:plyr':
##
##   arrange, mutate, rename, summarise

## The following object is masked from 'package:MASS':
##
##   select

## The following object is masked from 'package:stats':
##
##   filter
```

```

## The following object is masked from 'package:graphics':
##
##      layout

# Importing the Data
train <- read.table("D:\\Kaggle Projects\\Dengue\\dengue_features_train.csv",
header=TRUE, sep=",")
test <- read.table("D:\\Kaggle Projects\\Dengue\\dengue_features_test.csv",
header=TRUE, sep=",")

attach(train)
attach(test)

## The following objects are masked from train:
##
##      city, ndvi_ne, ndvi_nw, ndvi_se, ndvi_sw,
##      precipitation_amt_mm, reanalysis_air_temp_k,
##      reanalysis_avg_temp_k, reanalysis_dew_point_temp_k,
##      reanalysis_max_air_temp_k, reanalysis_min_air_temp_k,
##      reanalysis_precip_amt_kg_per_m2,
##      reanalysis_relative_humidity_percent,
##      reanalysis_sat_precip_amt_mm,
##      reanalysis_specific_humidity_g_per_kg, reanalysis_tdtr_k,
##      station_avg_temp_c, station_diur_temp_rng_c,
##      station_max_temp_c, station_min_temp_c, station_precip_mm,
##      week_start_date, weekofyear, year

#Exploring the Data
head(train,5)

##      city year weekofyear week_start_date  ndvi_ne  ndvi_nw  ndvi_se
## 1      1 1990          18      4/30/1990 0.1226000 0.1037250 0.1984833
## 2      1 1990          19      5/7/1990  0.1699000 0.1421750 0.1623571
## 3      1 1990          20      5/14/1990 0.0322500 0.1729667 0.1572000
## 4      1 1990          21      5/21/1990 0.1286333 0.2450667 0.2275571
## 5      1 1990          22      5/28/1990 0.1962000 0.2622000 0.2512000
##      ndvi_sw precipitation_amt_mm reanalysis_air_temp_k
## 1 0.1776167          12.42          297.5729
## 2 0.1554857          22.82          298.2114
## 3 0.1708429          34.54          298.7814
## 4 0.2358857          15.36          298.9871
## 5 0.2473400           7.52          299.5186
##      reanalysis_avg_temp_k reanalysis_dew_point_temp_k
## 1          297.7429          292.4143
## 2          298.4429          293.9514
## 3          298.8786          295.4343
## 4          299.2286          295.3100
## 5          299.6643          295.8214
##      reanalysis_max_air_temp_k reanalysis_min_air_temp_k
## 1          299.8          295.9
## 2          300.9          296.4

```

```
## 3          300.5          297.3
## 4          301.4          297.0
## 5          301.9          297.5
## reanalysis_precip_amt_kg_per_m2 reanalysis_relative_humidity_percent
## 1          32.00          73.36571
## 2          17.94          77.36857
## 3          26.10          82.05286
## 4          13.90          80.33714
## 5          12.20          80.46000
## reanalysis_sat_precip_amt_mm reanalysis_specific_humidity_g_per_kg
## 1          12.42          14.01286
## 2          22.82          15.37286
## 3          34.54          16.84857
## 4          15.36          16.67286
## 5          7.52          17.21000
## reanalysis_tdtr_k station_avg_temp_c station_diur_temp_rng_c
## 1          2.628571          25.44286          6.900000
## 2          2.371429          26.71429          6.371429
## 3          2.300000          26.71429          6.485714
## 4          2.428571          27.47143          6.771429
## 5          3.014286          28.94286          9.371429
## station_max_temp_c station_min_temp_c station_precip_mm total_cases
## 1          29.4          20.0          16.0          4
## 2          31.7          22.2          8.6          5
## 3          32.2          22.8          41.4          4
## 4          33.3          23.3          4.0          3
## 5          35.0          23.9          5.8          6
```

#getting the Data over the Time

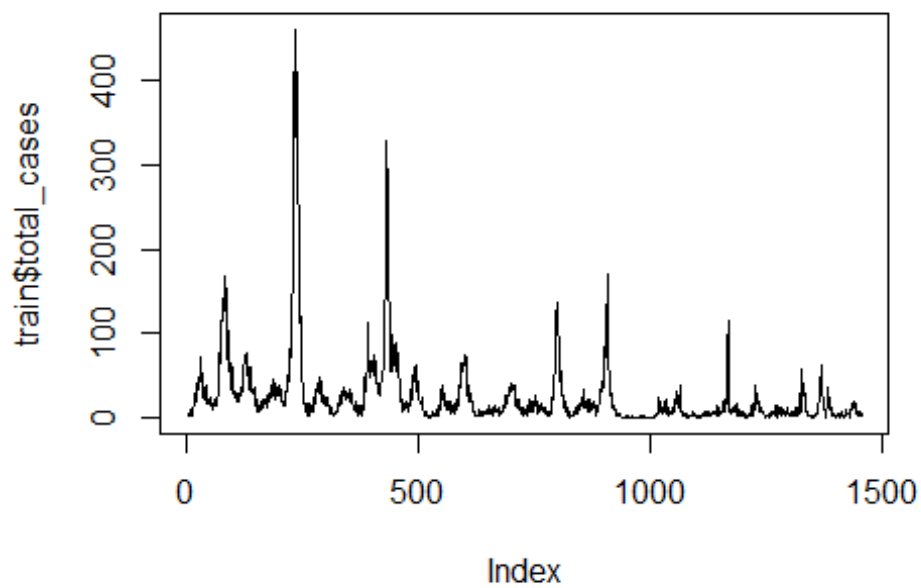
```
aggdata <- aggregate(train$city, by=list(train$year,train$city),FUN=mean,
na.rm=TRUE)
```

#Conclusion here is that the Sj has the Data from 1990 to 2008

#where as the iq has the Data from 2000 to 2010

#plotting the Data based on the Number of Cases

```
plot(train$total_cases,type="l")
```



```
plot(train$year,train$total_cases,type = "h" , col="red")
```



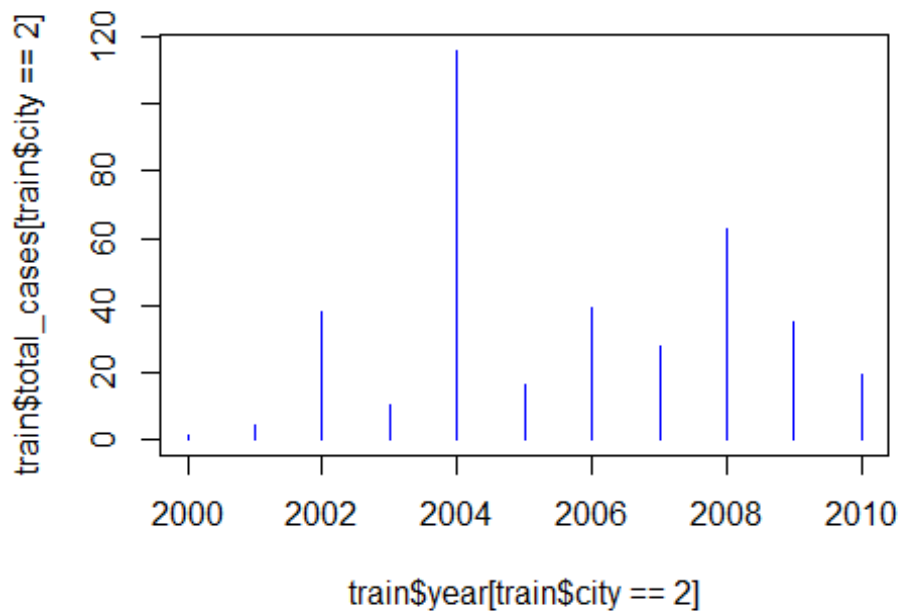
```
# As we can see the Cases are distributed over the year  
# But we dont know for which city it is for as we have two city
```

```
unique(train$city)
```

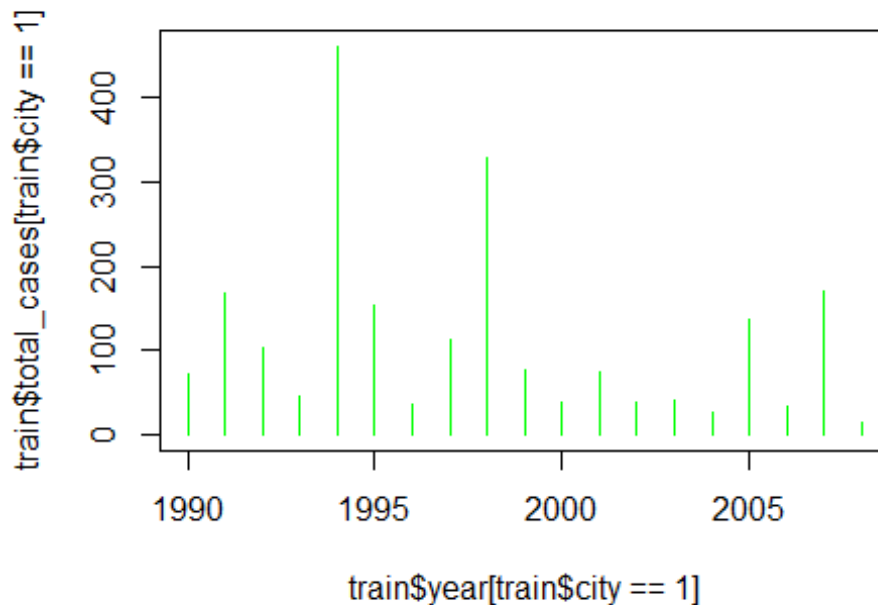
```
## [1] 1 2
```

```
#two cities are Sj = 1 and iq = 2
```

```
plot(train$year[train$city==2],train$total_cases[train$city==2],type="h" ,  
col="blue")
```



```
plot(train$year[train$city==1],train$total_cases[train$city==1],type="h" ,  
col="green")
```



```
# Conclusion
# For City Sj the Maximum number of cases are in 1994
# For City iq the Maximum number of cases are in 2004

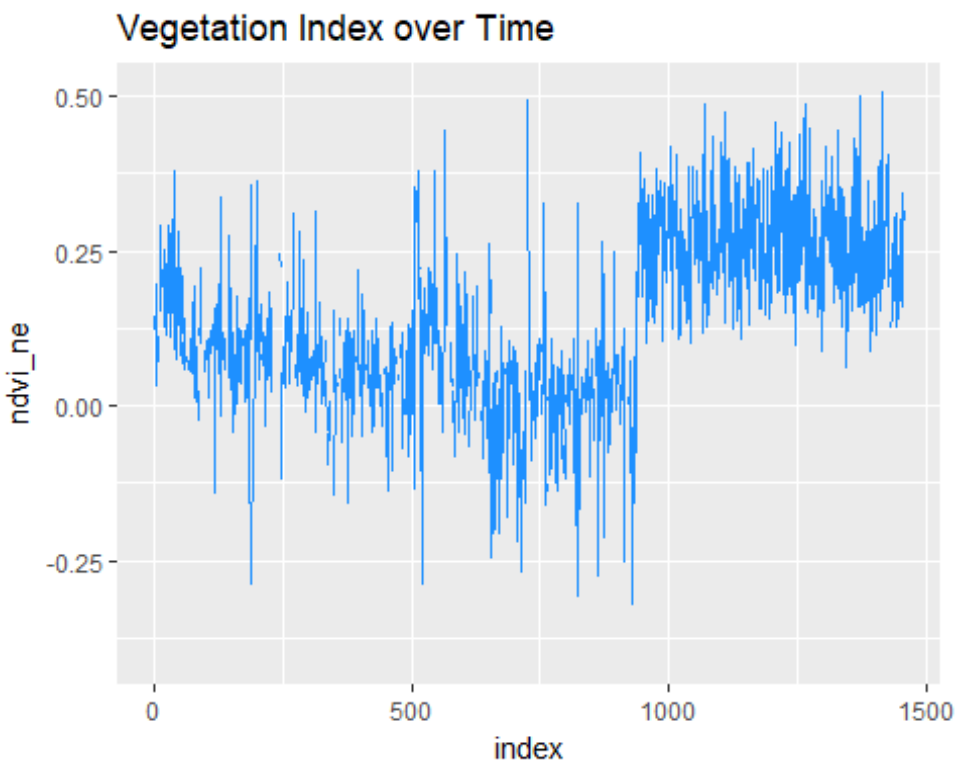
# count missing values (as percent)
apply(train, 2, function(x)
  round(100 * (length(which(is.na(x))))/length(x) , digits = 1)) %>%
  as.data.frame() %>%
  `names<-`('Percent of Missing Values')

##                                Percent of Missing Values
## city                                0.0
## year                                0.0
## weekofyear                          0.0
## week_start_date                     0.0
## ndvi_ne                             13.3
## ndvi_nw                             3.6
## ndvi_se                             1.5
## ndvi_sw                             1.5
## precipitation_amt_mm                 0.9
## reanalysis_air_temp_k                0.7
## reanalysis_avg_temp_k                0.7
## reanalysis_dew_point_temp_k          0.7
## reanalysis_max_air_temp_k            0.7
## reanalysis_min_air_temp_k            0.7
## reanalysis_precip_amt_kg_per_m2      0.7
## reanalysis_relative_humidity_percent  0.7
```

```
## reanalysis_sat_precip_amt_mm 0.9
## reanalysis_specific_humidity_g_per_kg 0.7
## reanalysis_tdtr_k 0.7
## station_avg_temp_c 3.0
## station_diur_temp_rng_c 3.0
## station_max_temp_c 1.4
## station_min_temp_c 1.0
## station_precip_mm 1.5
## total_cases 0.0
```

Plotting the Data

```
train %>%
  mutate(index = as.numeric(row.names(.))) %>%
  ggplot(aes(index, ndvi_ne)) +
  geom_line(colour = 'dodgerblue') +
  ggtitle("Vegetation Index over Time")
```

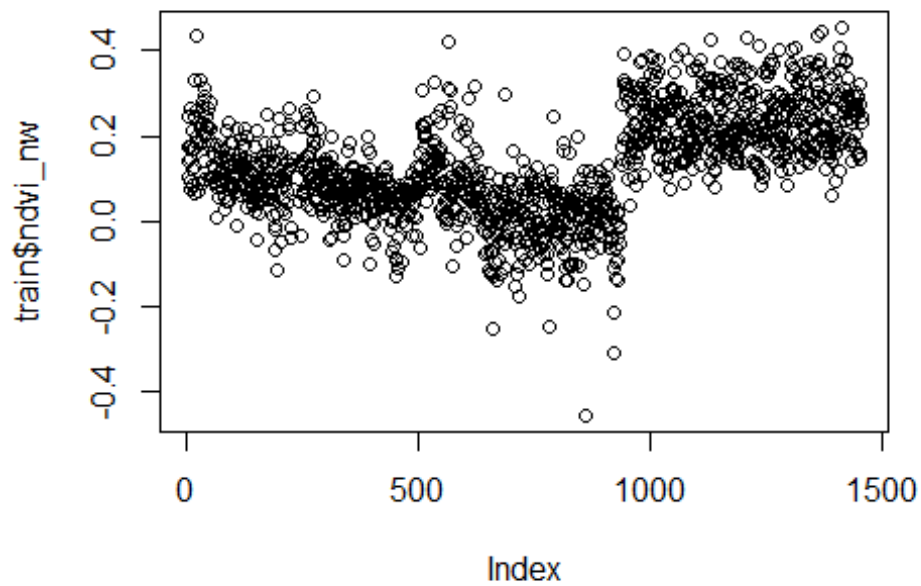


Dropping the Coloum with 13% of the missing Data

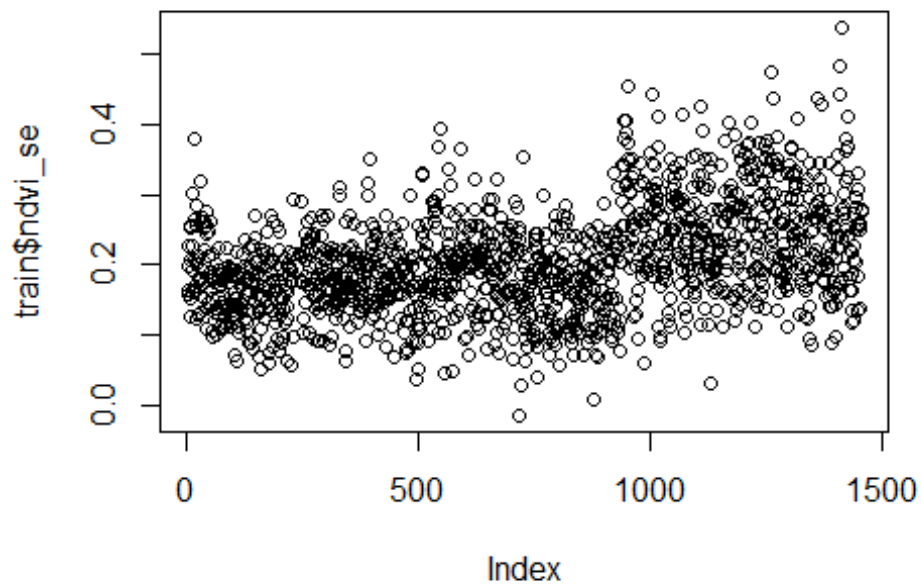
```
train$ndvi_ne <- NULL
test$ndvi_ne <- NULL
```

Replacing the Values with the Mean

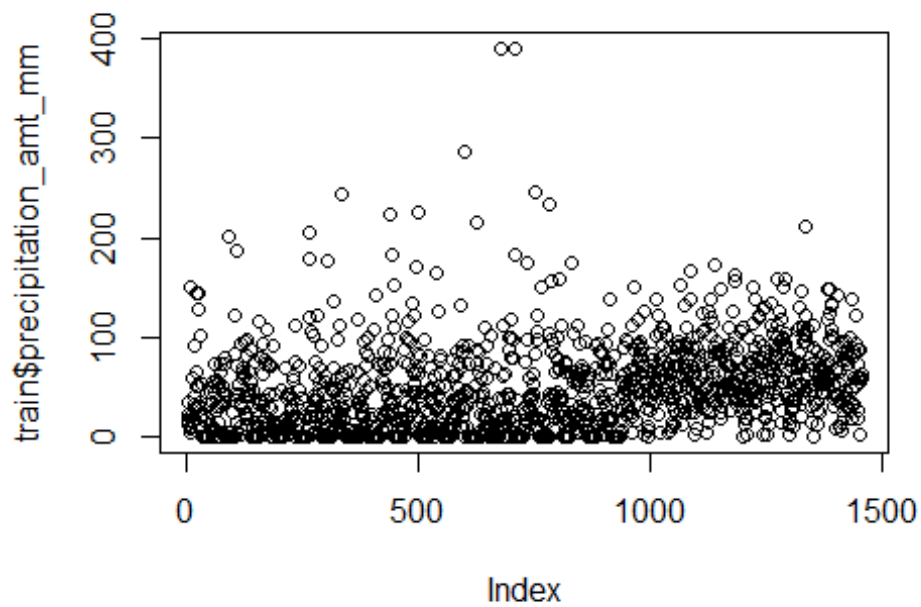
```
plot(train$ndvi_nw)
```

```
mean(train$ndvi_nw, na.rm = TRUE)
## [1] 0.1305526
train$ndvi_nw[is.na(train$ndvi_nw)] <- mean(train$ndvi_nw, na.rm = TRUE)
plot(train$ndvi_se)
```



```
mean(train$ndvi_se, na.rm = TRUE)
## [1] 0.2037832
train$ndvi_se[is.na(train$ndvi_se)] <- mean(train$ndvi_se, na.rm = TRUE)
plot(train$precipitation_amt_mm)
```

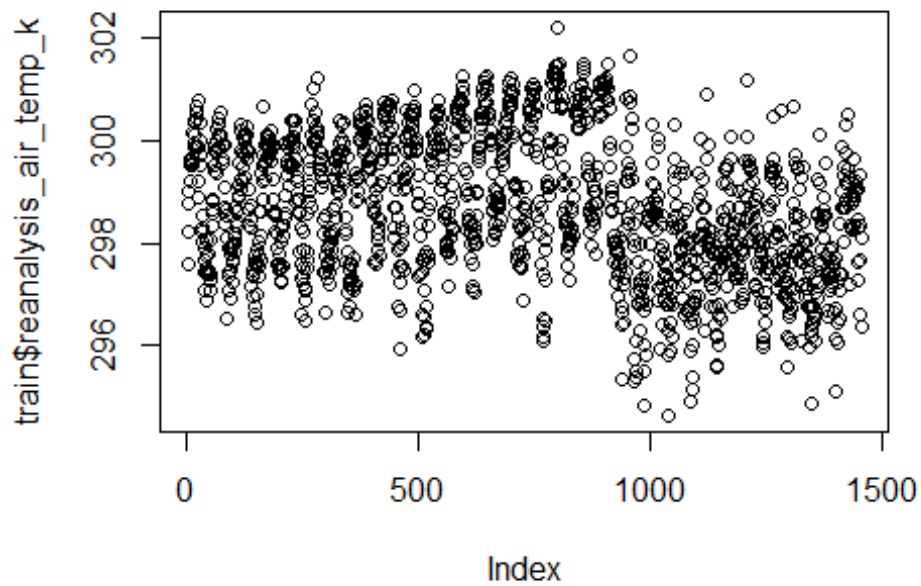


```
mean(train$precipitation_amt_mm,na.rm = TRUE)

## [1] 45.76039

train$precipitation_amt_mm[is.na(train$precipitation_amt_mm)] <-
mean(train$precipitation_amt_mm,na.rm = TRUE)

plot(train$reanalysis_air_temp_k)
```

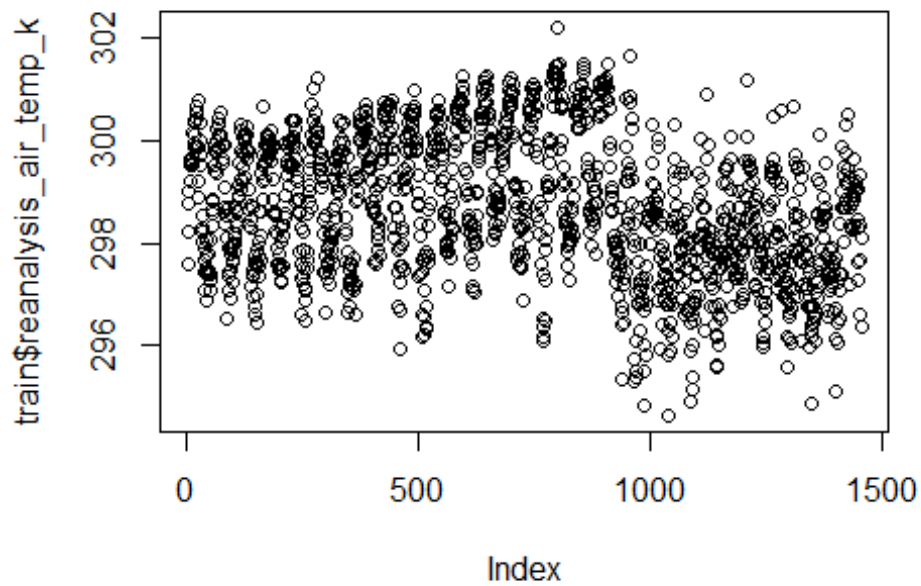


```
mean(train$reanalysis_air_temp_k, na.rm = TRUE)

## [1] 298.7019

train$reanalysis_air_temp_k[is.na(train$reanalysis_air_temp_k)] <-
mean(train$reanalysis_air_temp_k, na.rm = TRUE)

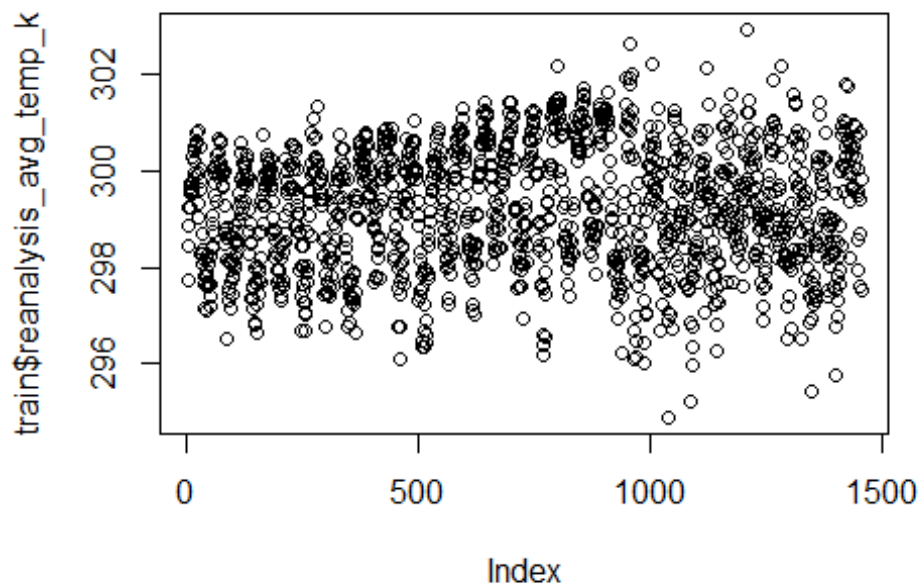
plot(train$reanalysis_air_temp_k)
```



```
mean(train$reanalysis_air_temp_k, na.rm = TRUE)
## [1] 298.7019

train$reanalysis_air_temp_k[is.na(train$reanalysis_air_temp_k)] <-
mean(train$reanalysis_air_temp_k, na.rm = TRUE)

plot(train$reanalysis_avg_temp_k)
```

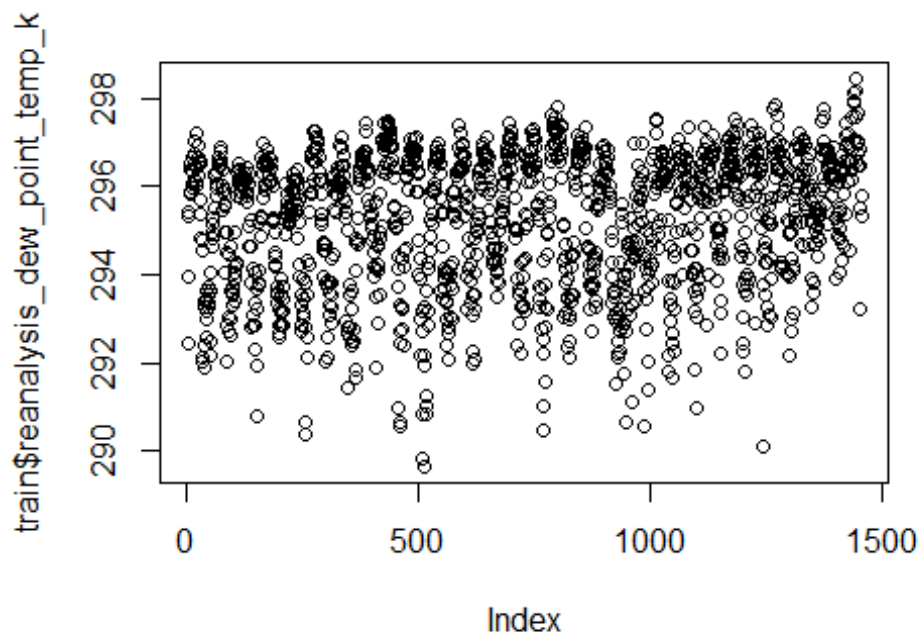


```
mean(train$reanalysis_avg_temp_k, na.rm = TRUE)

## [1] 299.2256

train$reanalysis_avg_temp_k[is.na(train$reanalysis_avg_temp_k)] <-
mean(train$reanalysis_avg_temp_k, na.rm = TRUE)

plot(train$reanalysis_dew_point_temp_k)
```

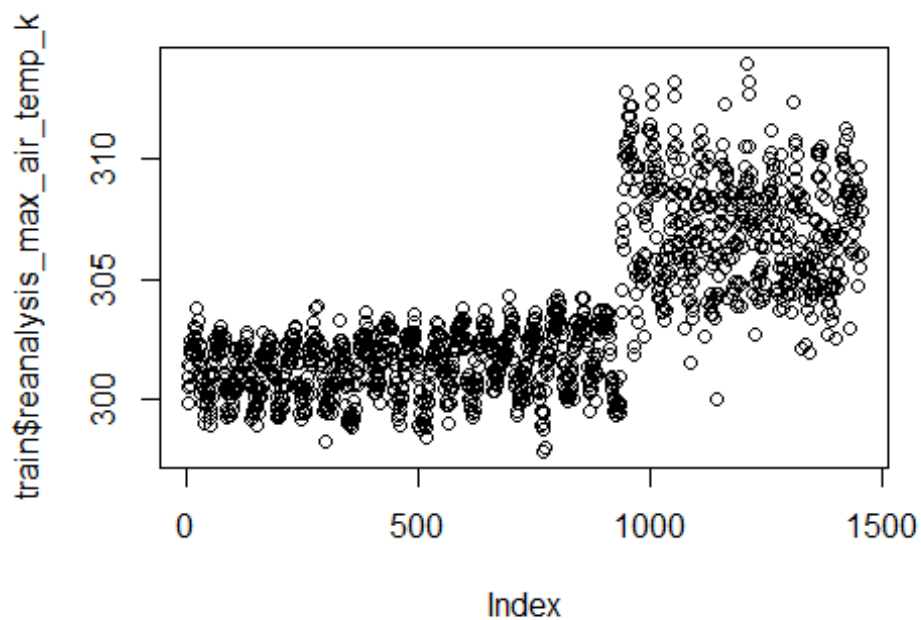


```
mean(train$reanalysis_dew_point_temp_k, na.rm = TRUE)

## [1] 295.2464

train$reanalysis_dew_point_temp_k[is.na(train$reanalysis_dew_point_temp_k)]
<- mean(train$reanalysis_dew_point_temp_k, na.rm = TRUE)

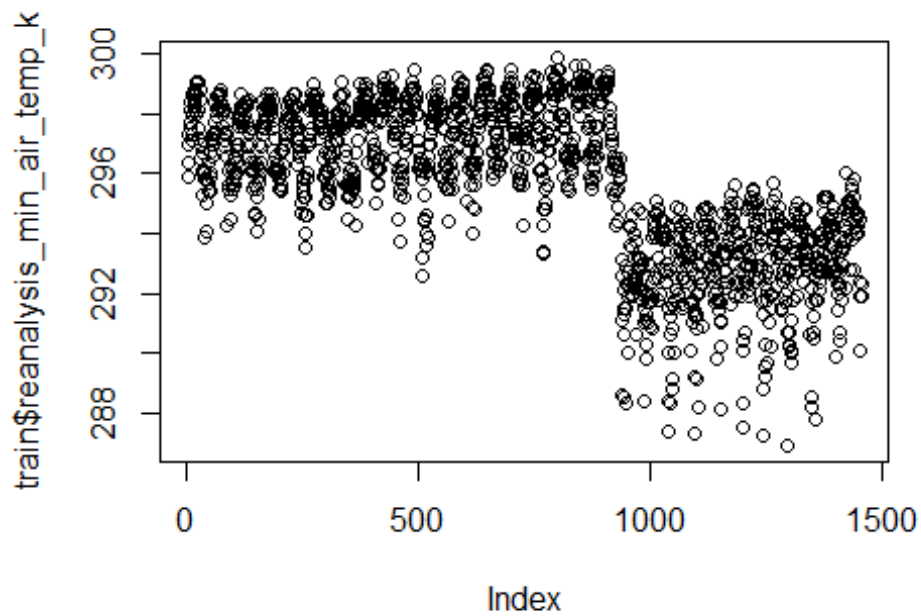
plot(train$reanalysis_max_air_temp_k)
```



```
mean(train$reanalysis_max_air_temp_k, na.rm = TRUE)
## [1] 303.4271

train$reanalysis_max_air_temp_k[is.na(train$reanalysis_max_air_temp_k)] <-
mean(train$reanalysis_max_air_temp_k, na.rm = TRUE)

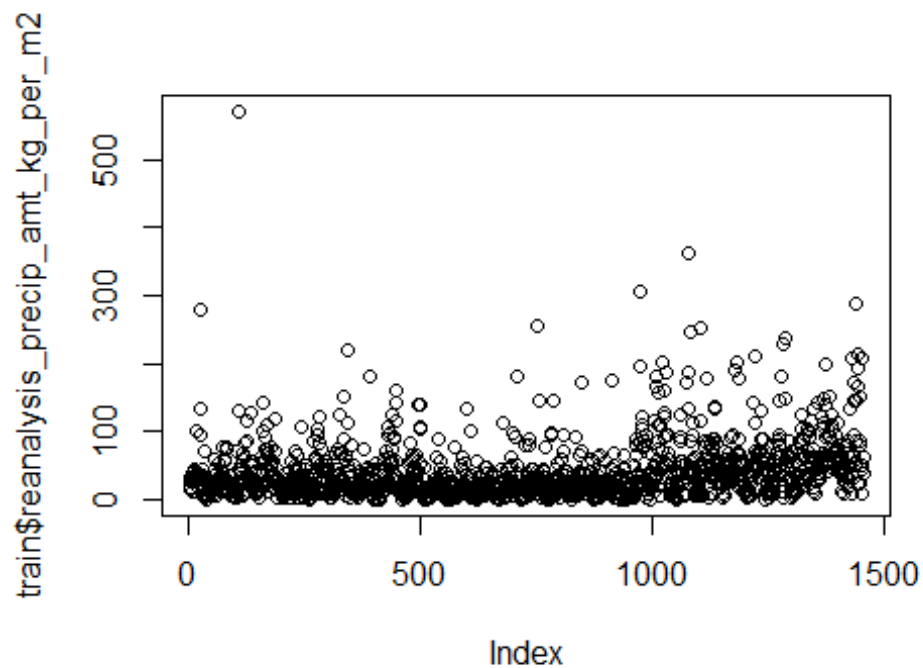
plot(train$reanalysis_min_air_temp_k)
```

```
mean(train$reanalysis_min_air_temp_k, na.rm = TRUE)
## [1] 295.7192

train$reanalysis_min_air_temp_k[is.na(train$reanalysis_min_air_temp_k)] <-
mean(train$reanalysis_min_air_temp_k, na.rm = TRUE)

plot(train$reanalysis_precip_amt_kg_per_m2)
```

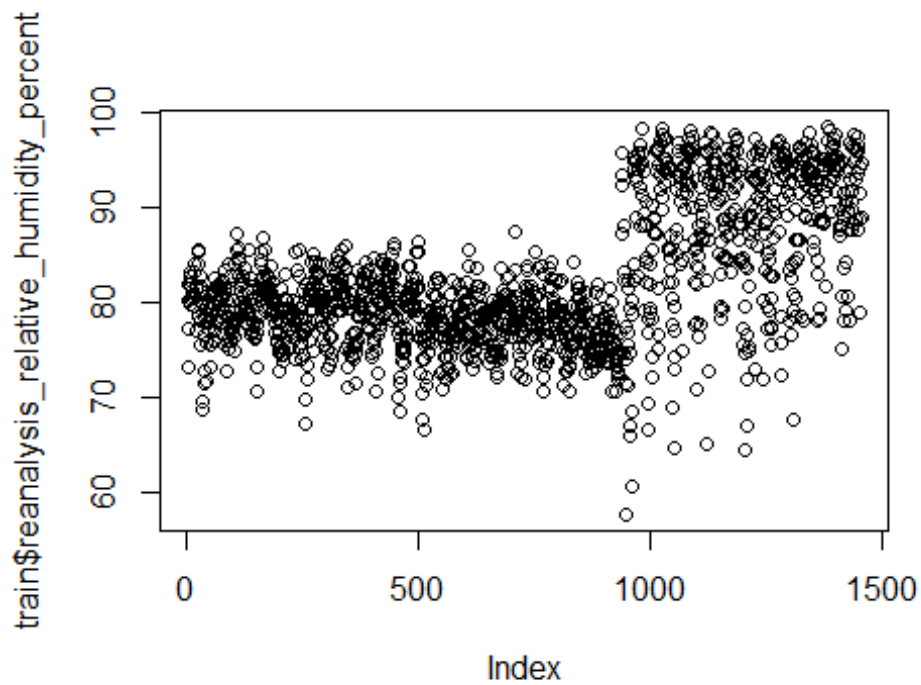


```
mean(train$reanalysis_precip_amt_kg_per_m2, na.rm = TRUE)

## [1] 40.15182

train$reanalysis_precip_amt_kg_per_m2[is.na(train$reanalysis_precip_amt_kg_per_m2)] <- mean(train$reanalysis_precip_amt_kg_per_m2, na.rm = TRUE)

plot(train$reanalysis_relative_humidity_percent)
```

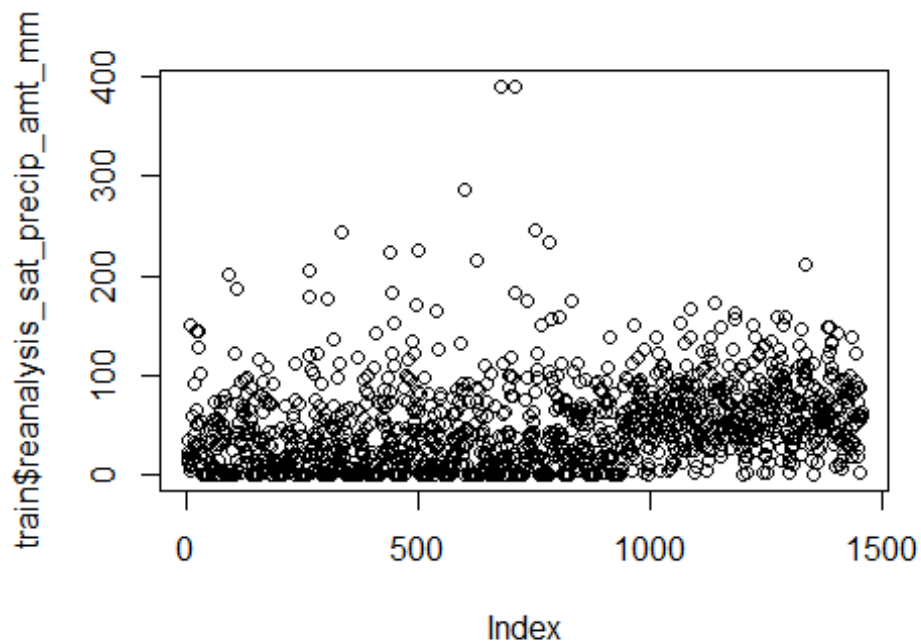


```
mean(train$reanalysis_relative_humidity_percent, na.rm = TRUE)
```

```
## [1] 82.16196
```

```
train$reanalysis_relative_humidity_percent[is.na(train$reanalysis_relative_humidity_percent)] <- mean(train$reanalysis_relative_humidity_percent, na.rm = TRUE)
```

```
plot(train$reanalysis_sat_precip_amt_mm)
```

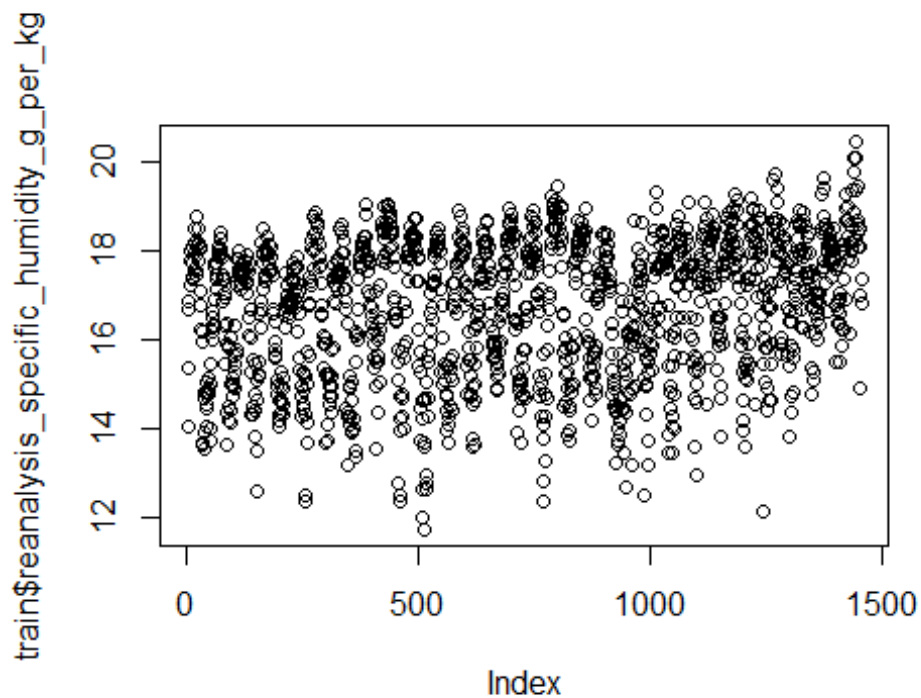


```
mean(train$reanalysis_sat_precip_amt_mm,na.rm = TRUE)

## [1] 45.76039

train$reanalysis_sat_precip_amt_mm[is.na(train$reanalysis_sat_precip_amt_mm)]
<- mean(train$reanalysis_sat_precip_amt_mm,na.rm = TRUE)

plot(train$reanalysis_specific_humidity_g_per_kg)
```

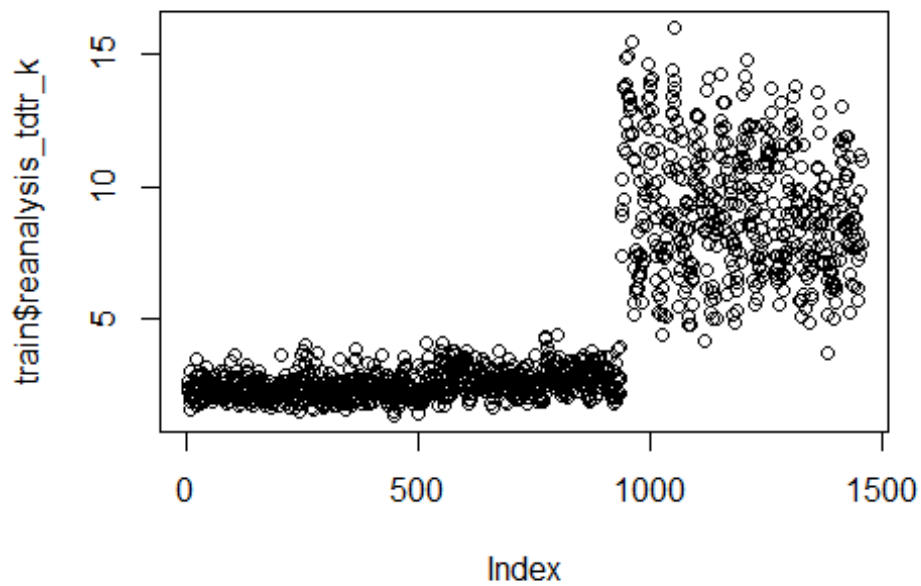


```
mean(train$reanalysis_specific_humidity_g_per_kg,na.rm = TRUE)
```

```
## [1] 16.74643
```

```
train$reanalysis_specific_humidity_g_per_kg[is.na(train$reanalysis_specific_humidity_g_per_kg)] <- mean(train$reanalysis_specific_humidity_g_per_kg,na.rm = TRUE)
```

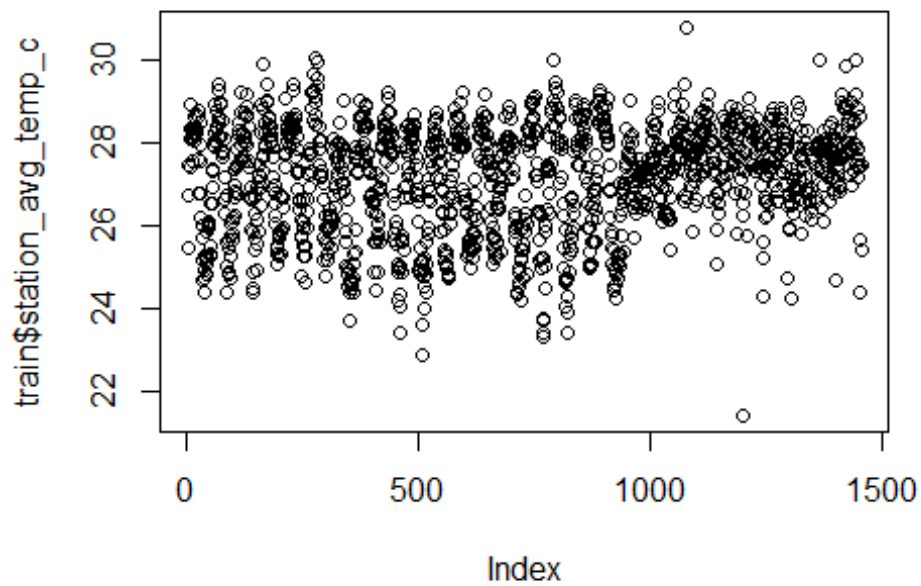
```
plot(train$reanalysis_tdtr_k)
```



```
mean(train$reanalysis_tdtr_k, na.rm = TRUE)
## [1] 4.903754

train$reanalysis_tdtr_k[is.na(train$reanalysis_tdtr_k)] <-
mean(train$reanalysis_tdtr_k, na.rm = TRUE)

plot(train$station_avg_temp_c)
```

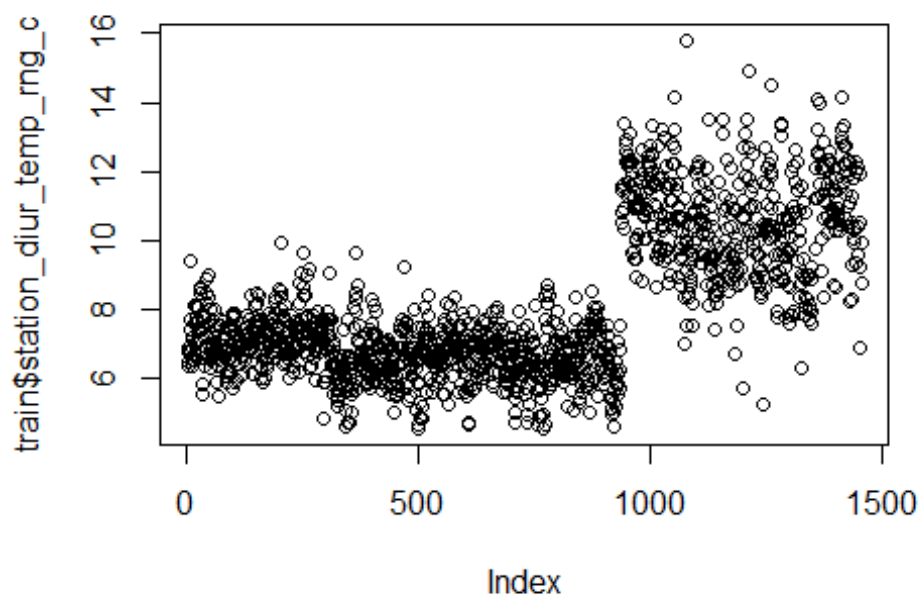


```
mean(train$station_avg_temp_c, na.rm = TRUE)

## [1] 27.18578

train$station_avg_temp_c[is.na(train$station_avg_temp_c)] <-
mean(train$station_avg_temp_c, na.rm = TRUE)

plot(train$station_diur_temp_rng_c)
```

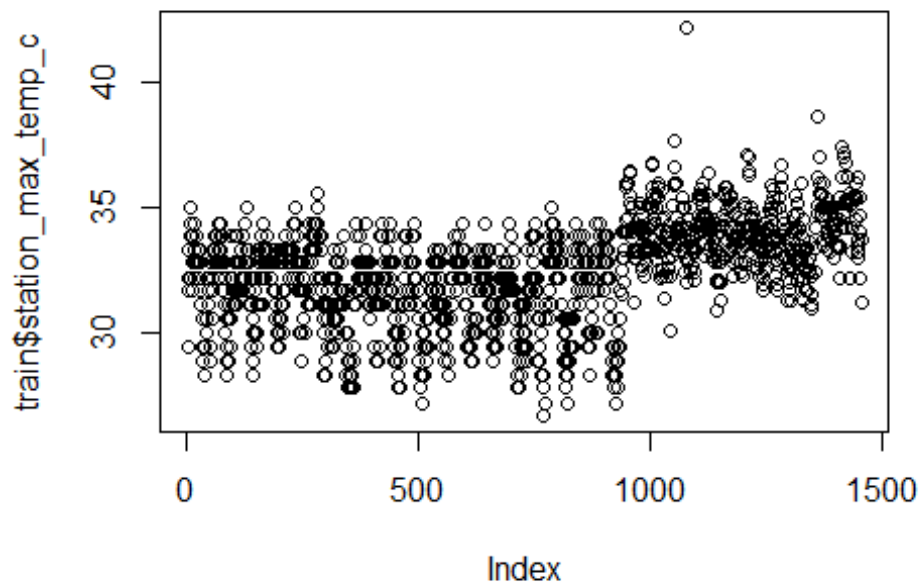


```
mean(train$station_diur_temp_rng_c, na.rm = TRUE)

## [1] 8.059328

train$station_diur_temp_rng_c[is.na(train$station_diur_temp_rng_c)] <-
mean(train$station_diur_temp_rng_c, na.rm = TRUE)

plot(train$station_max_temp_c)
```

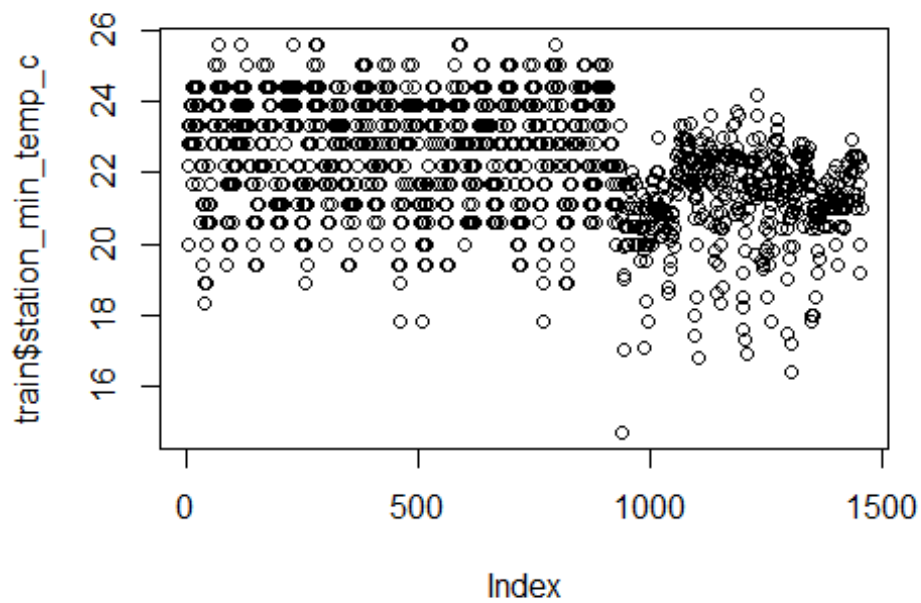



```
mean(train$station_max_temp_c, na.rm = TRUE)

## [1] 32.45244

train$station_max_temp_c[is.na(train$station_max_temp_c)] <-
mean(train$station_max_temp_c, na.rm = TRUE)

plot(train$station_min_temp_c)
```

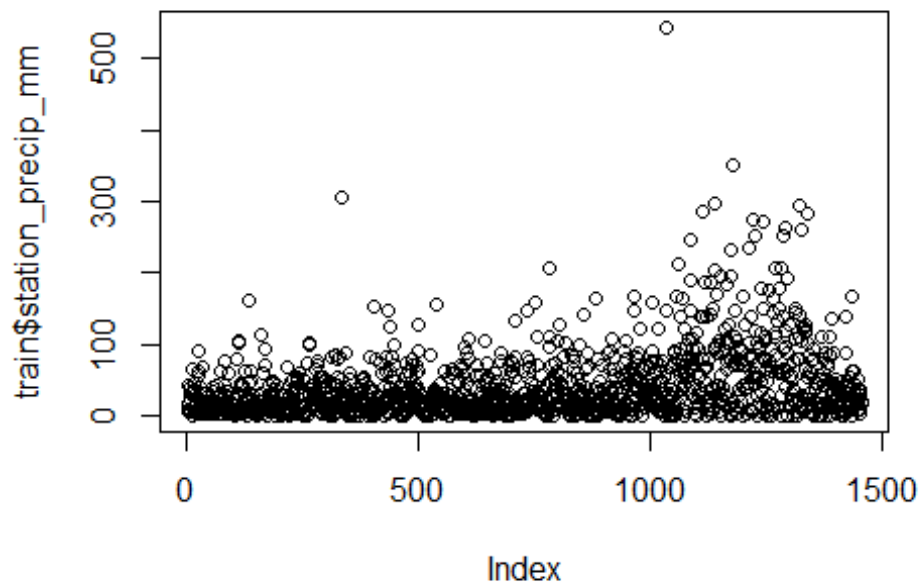


```
mean(train$station_min_temp_c, na.rm = TRUE)

## [1] 22.10215

train$station_min_temp_c[is.na(train$station_min_temp_c)] <-
mean(train$station_min_temp_c, na.rm = TRUE)

plot(train$station_precip_mm)
```



```
mean(train$station_precip_mm, na.rm = TRUE)

## [1] 39.32636

train$station_precip_mm[is.na(train$station_precip_mm)] <-
mean(train$station_precip_mm, na.rm = TRUE)

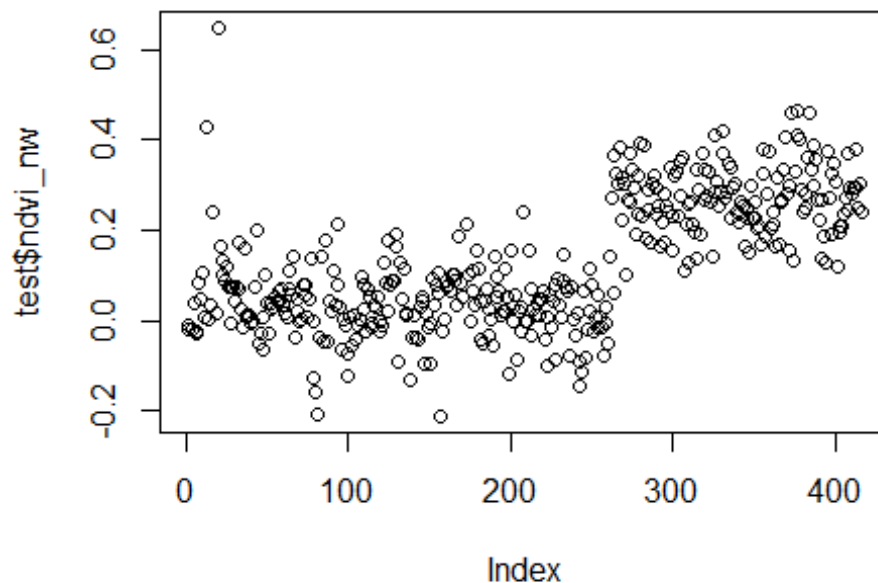
# count missing values (as percent)
apply(train, 2, function(x)
  round(100 * (length(which(is.na(x))))/length(x) , digits = 1)) %>%
  as.data.frame() %>%
  `names<-`('Percent of Missing Values')

##                               Percent of Missing Values
## city                               0.0
## year                               0.0
## weekofyear                         0.0
## week_start_date                   0.0
## ndvi_nw                           0.0
## ndvi_se                           0.0
## ndvi_sw                           1.5
## precipitation_amt_mm              0.0
## reanalysis_air_temp_k             0.0
## reanalysis_avg_temp_k             0.0
## reanalysis_dew_point_temp_k       0.0
## reanalysis_max_air_temp_k         0.0
```

```
## reanalysis_min_air_temp_k 0.0
## reanalysis_precip_amt_kg_per_m2 0.0
## reanalysis_relative_humidity_percent 0.0
## reanalysis_sat_precip_amt_mm 0.0
## reanalysis_specific_humidity_g_per_kg 0.0
## reanalysis_tdtr_k 0.0
## station_avg_temp_c 0.0
## station_diur_temp_rng_c 0.0
## station_max_temp_c 0.0
## station_min_temp_c 0.0
## station_precip_mm 0.0
## total_cases 0.0
```

Replacing the Values with the Mean of Test

```
plot(test$ndvi_nw)
```

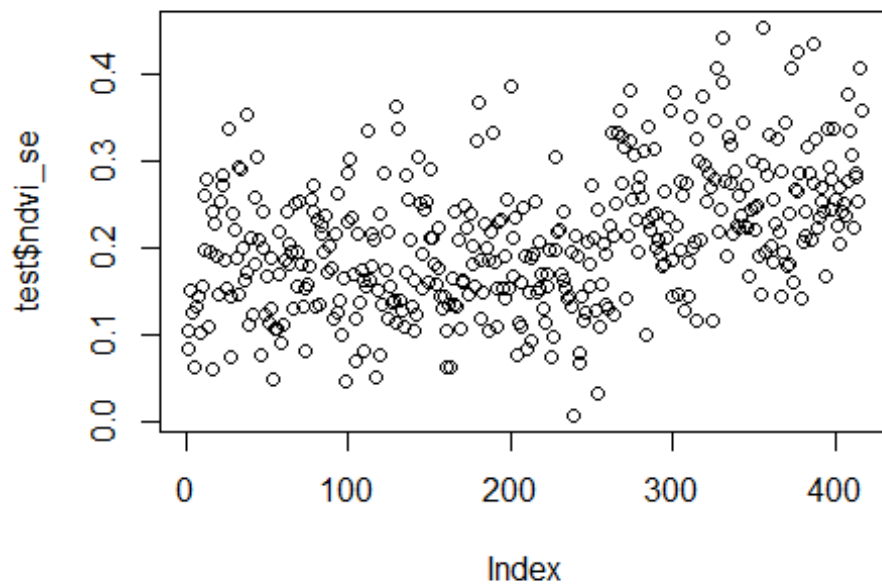


```
mean(test$ndvi_nw,na.rm = TRUE)
```

```
## [1] 0.126803
```

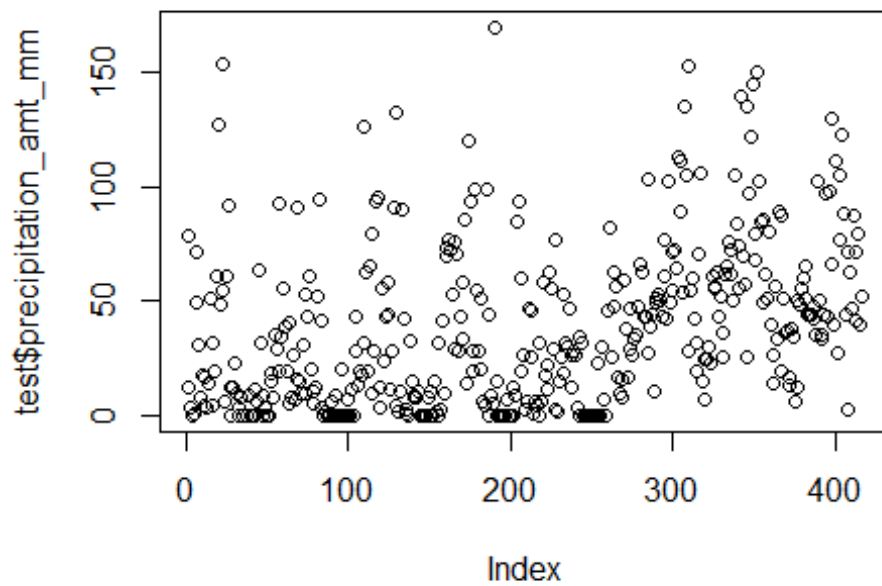
```
test$ndvi_nw[is.na(test$ndvi_nw)] <- mean(test$ndvi_nw,na.rm = TRUE)
```

```
plot(test$ndvi_se)
```



```
mean(test$ndvi_se, na.rm = TRUE)
## [1] 0.2077017
test$ndvi_se[is.na(test$ndvi_se)] <- mean(test$ndvi_se, na.rm = TRUE)

plot(test$precipitation_amt_mm)
```

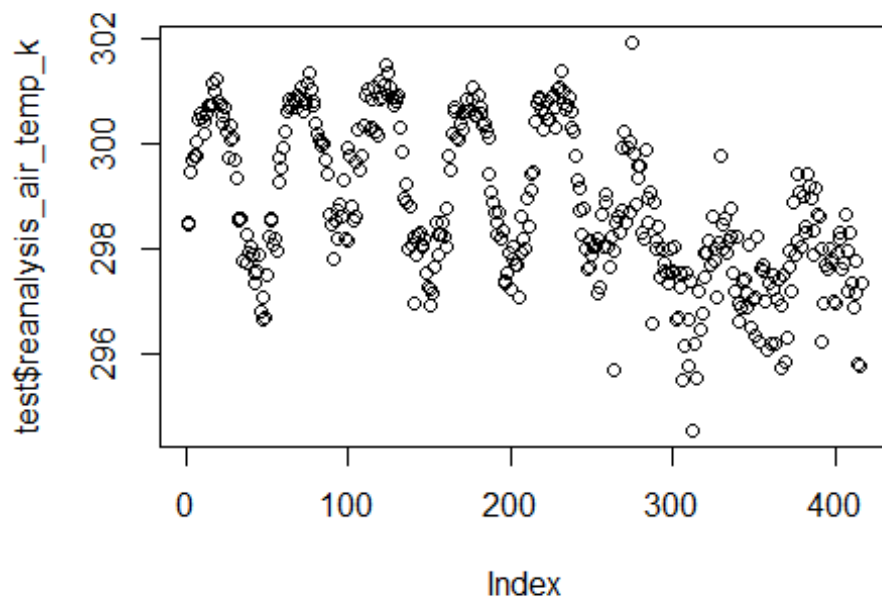


```
mean(test$precipitation_amt_mm, na.rm = TRUE)

## [1] 38.35432

test$precipitation_amt_mm[is.na(test$precipitation_amt_mm)] <-
mean(test$precipitation_amt_mm, na.rm = TRUE)

plot(test$reanalysis_air_temp_k)
```

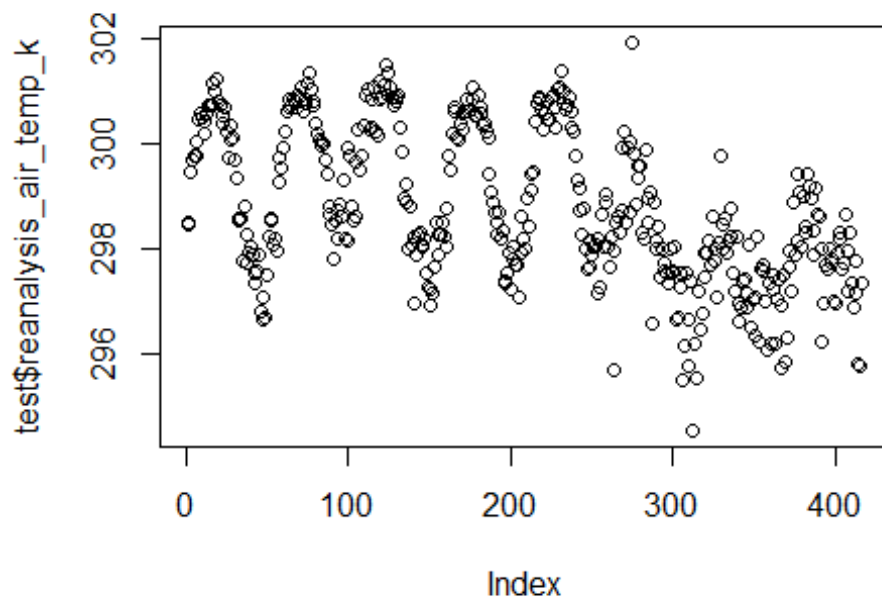


```
mean(test$reanalysis_air_temp_k, na.rm = TRUE)

## [1] 298.8183

test$reanalysis_air_temp_k[is.na(test$reanalysis_air_temp_k)] <-
mean(test$reanalysis_air_temp_k, na.rm = TRUE)

plot(test$reanalysis_air_temp_k)
```

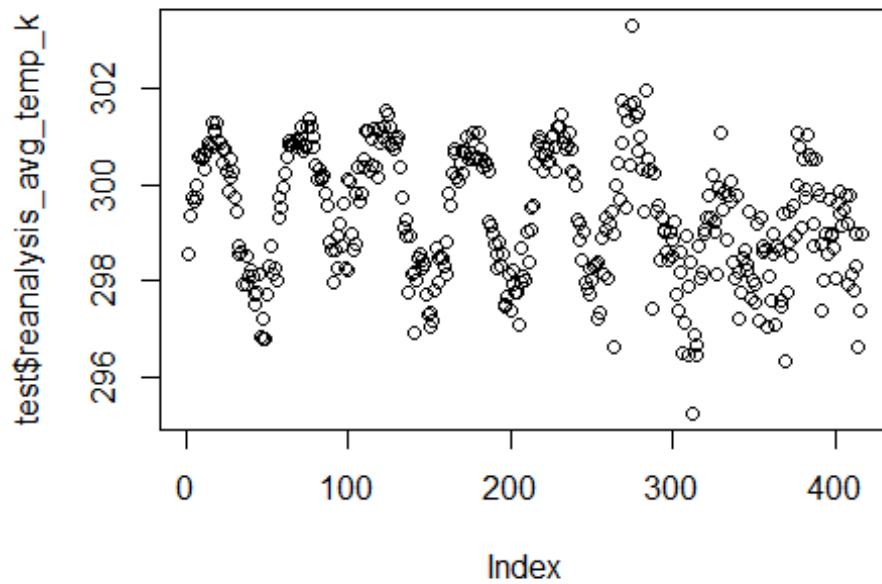


```
mean(test$reanalysis_air_temp_k, na.rm = TRUE)

## [1] 298.8183

test$reanalysis_air_temp_k[is.na(test$reanalysis_air_temp_k)] <-
mean(test$reanalysis_air_temp_k, na.rm = TRUE)

plot(test$reanalysis_avg_temp_k)
```

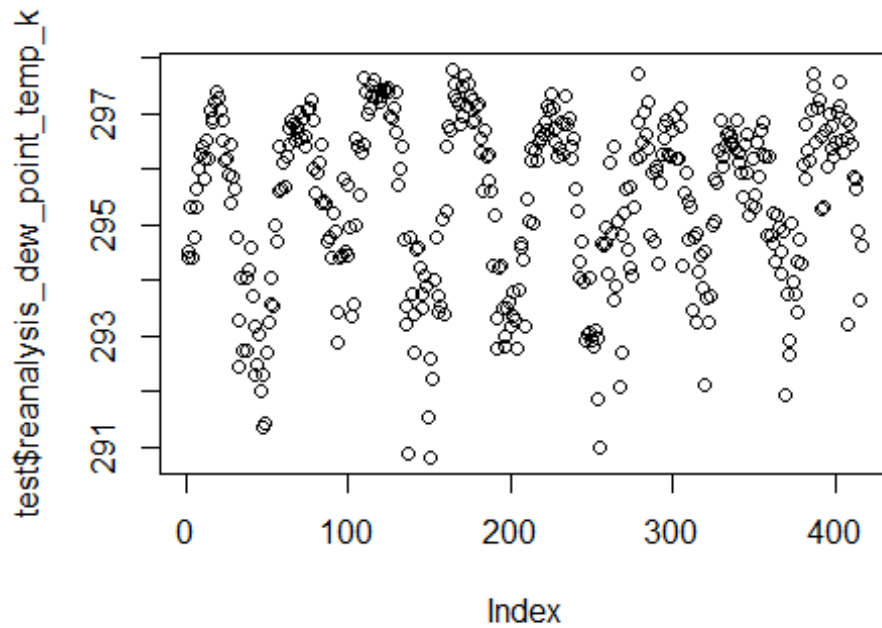



```
mean(test$reanalysis_avg_temp_k, na.rm = TRUE)

## [1] 299.3531

test$reanalysis_avg_temp_k[is.na(test$reanalysis_avg_temp_k)] <-
mean(test$reanalysis_avg_temp_k, na.rm = TRUE)

plot(test$reanalysis_dew_point_temp_k)
```

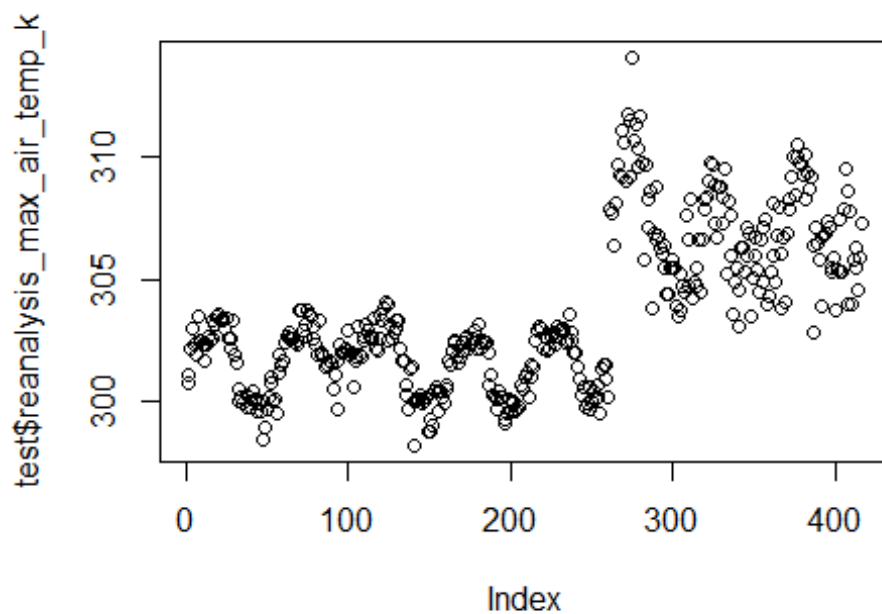


```
mean(test$reanalysis_dew_point_temp_k, na.rm = TRUE)

## [1] 295.4192

test$reanalysis_dew_point_temp_k[is.na(test$reanalysis_dew_point_temp_k)] <-
mean(test$reanalysis_dew_point_temp_k, na.rm = TRUE)

plot(test$reanalysis_max_air_temp_k)
```

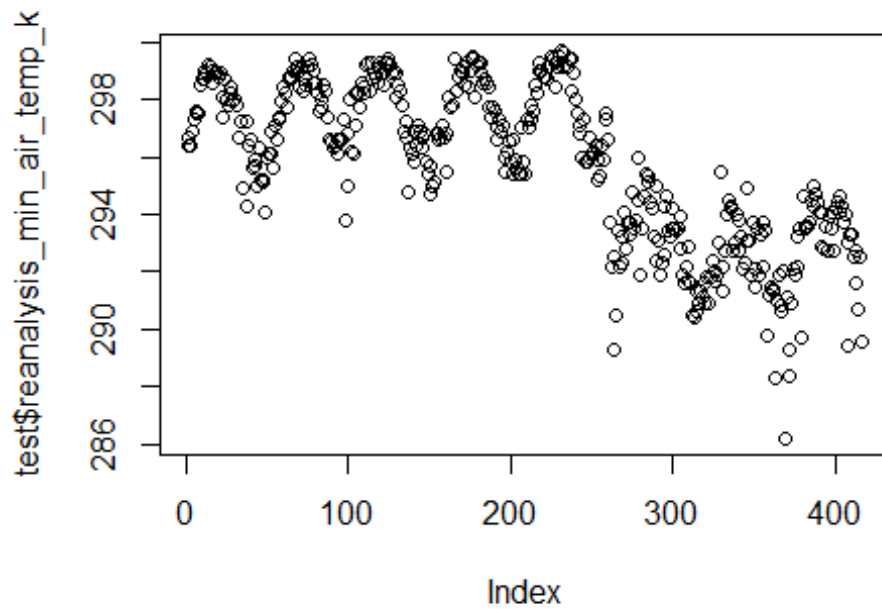


```
mean(test$reanalysis_max_air_temp_k, na.rm = TRUE)

## [1] 303.6234

test$reanalysis_max_air_temp_k[is.na(test$reanalysis_max_air_temp_k)] <-
mean(test$reanalysis_max_air_temp_k, na.rm = TRUE)

plot(test$reanalysis_min_air_temp_k)
```

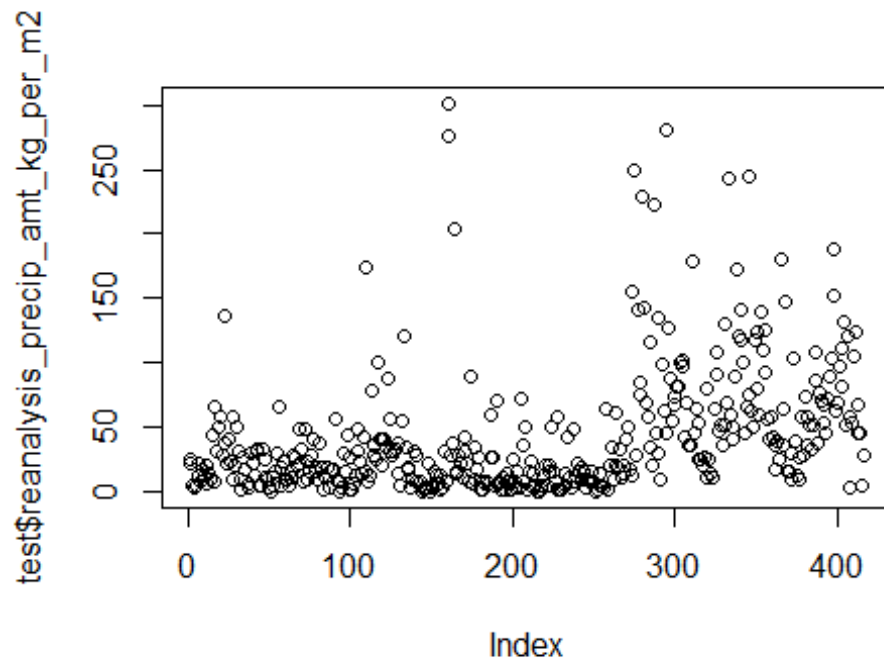


```
mean(test$reanalysis_min_air_temp_k, na.rm = TRUE)

## [1] 295.7435

test$reanalysis_min_air_temp_k[is.na(test$reanalysis_min_air_temp_k)] <-
mean(test$reanalysis_min_air_temp_k, na.rm = TRUE)

plot(test$reanalysis_precip_amt_kg_per_m2)
```

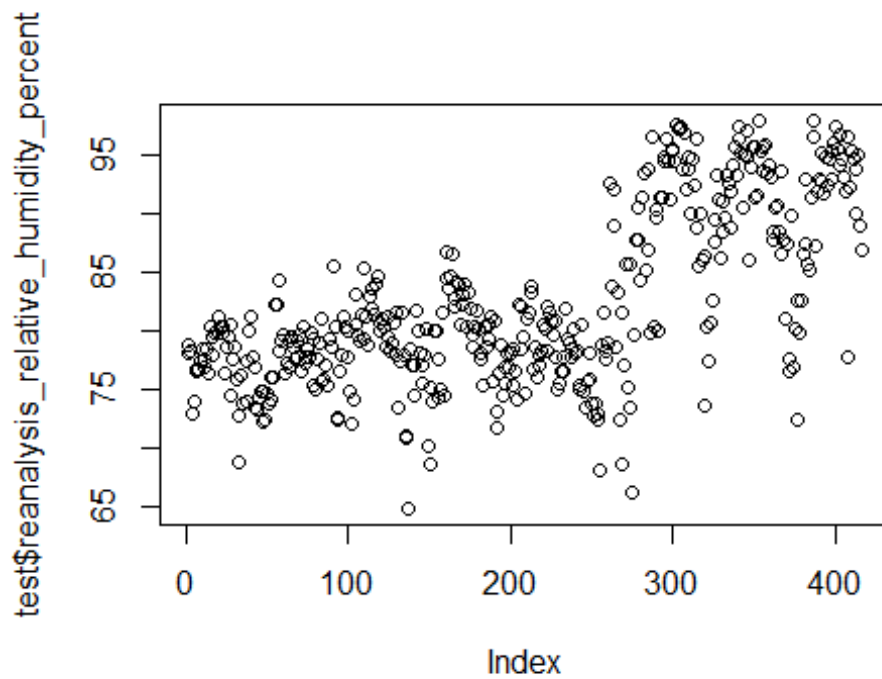


```
mean(test$reanalysis_precip_amt_kg_per_m2, na.rm = TRUE)
```

```
## [1] 42.17114
```

```
test$reanalysis_precip_amt_kg_per_m2[is.na(test$reanalysis_precip_amt_kg_per_m2)] <- mean(test$reanalysis_precip_amt_kg_per_m2, na.rm = TRUE)
```

```
plot(test$reanalysis_relative_humidity_percent)
```

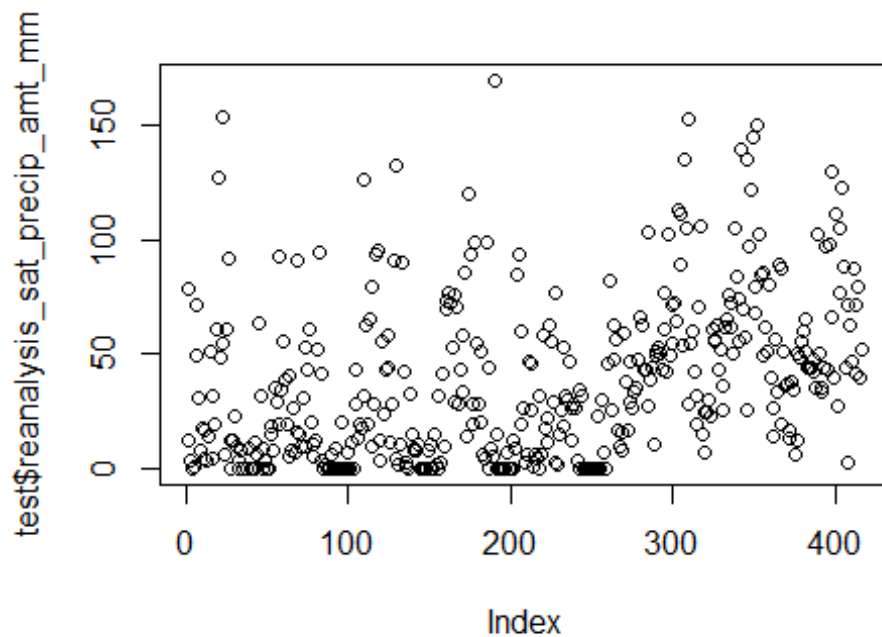


```
mean(test$reanalysis_relative_humidity_percent, na.rm = TRUE)
```

```
## [1] 82.49981
```

```
test$reanalysis_relative_humidity_percent[is.na(test$reanalysis_relative_humidity_percent)] <- mean(test$reanalysis_relative_humidity_percent, na.rm = TRUE)
```

```
plot(test$reanalysis_sat_precip_amt_mm)
```

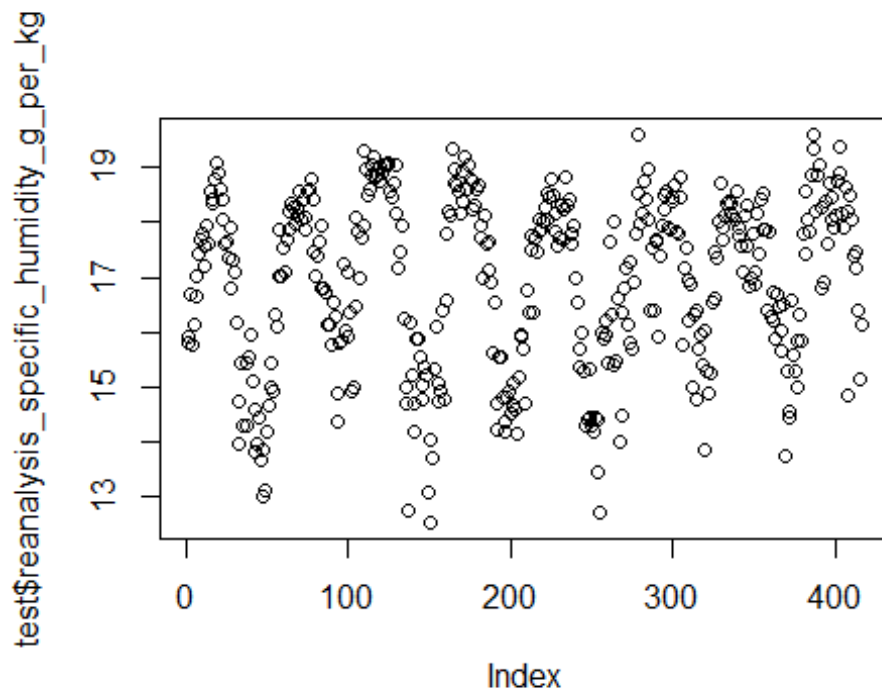


```
mean(test$reanalysis_sat_precip_amt_mm,na.rm = TRUE)
```

```
## [1] 38.35432
```

```
test$reanalysis_sat_precip_amt_mm[is.na(test$reanalysis_sat_precip_amt_mm)]  
<- mean(test$reanalysis_sat_precip_amt_mm,na.rm = TRUE)
```

```
plot(test$reanalysis_specific_humidity_g_per_kg)
```

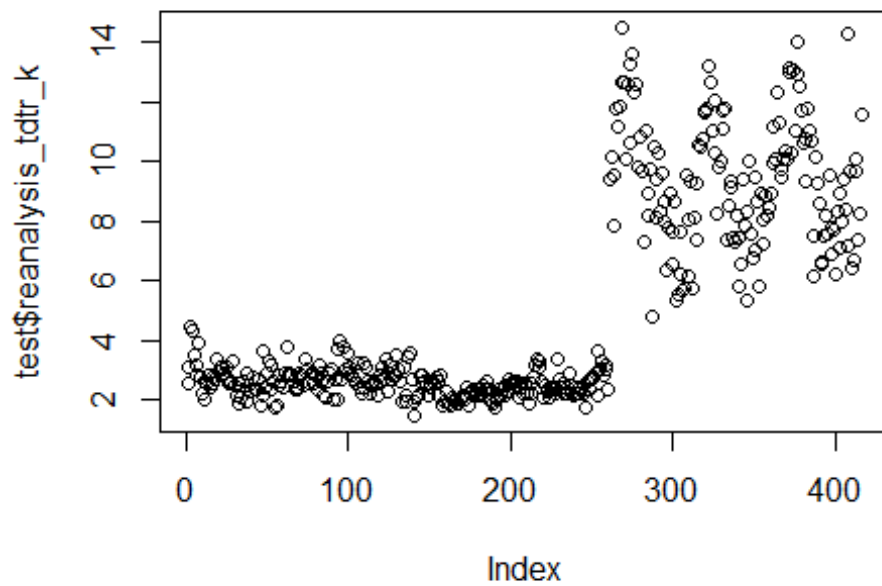


```
mean(test$reanalysis_specific_humidity_g_per_kg, na.rm = TRUE)

## [1] 16.92709

test$reanalysis_specific_humidity_g_per_kg[is.na(test$reanalysis_specific_hum
idity_g_per_kg)] <- mean(test$reanalysis_specific_humidity_g_per_kg, na.rm =
TRUE)

plot(test$reanalysis_tdtr_k)
```

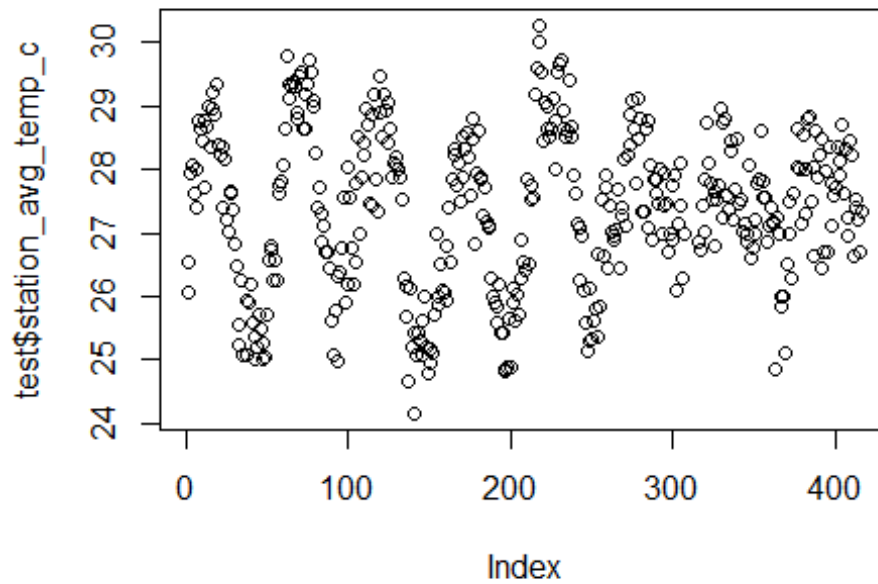



```
mean(test$reanalysis_tdtr_k, na.rm = TRUE)

## [1] 5.124569

test$reanalysis_tdtr_k[is.na(test$reanalysis_tdtr_k)] <-
mean(test$reanalysis_tdtr_k, na.rm = TRUE)

plot(test$station_avg_temp_c)
```

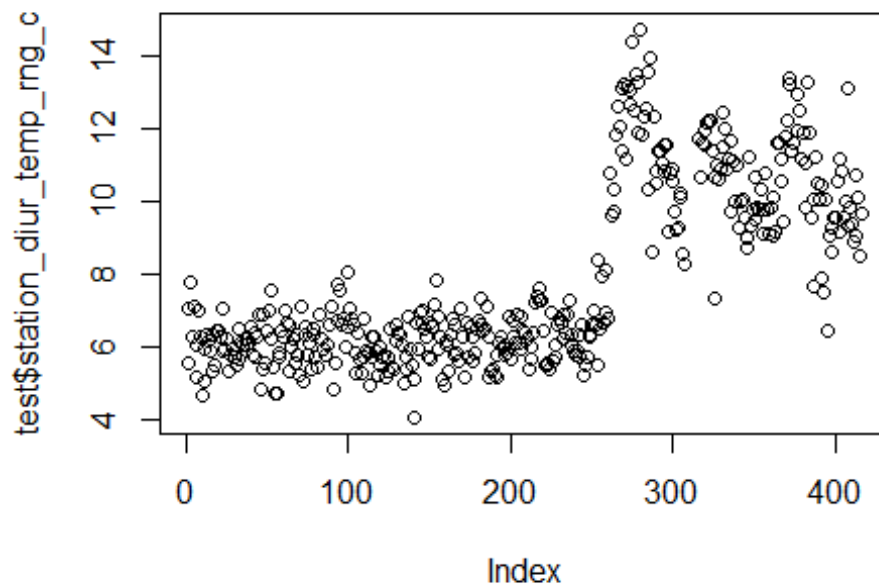


```
mean(test$station_avg_temp_c, na.rm = TRUE)

## [1] 27.36959

test$station_avg_temp_c[is.na(test$station_avg_temp_c)] <-
mean(test$station_avg_temp_c, na.rm = TRUE)

plot(test$station_diur_temp_rng_c)
```

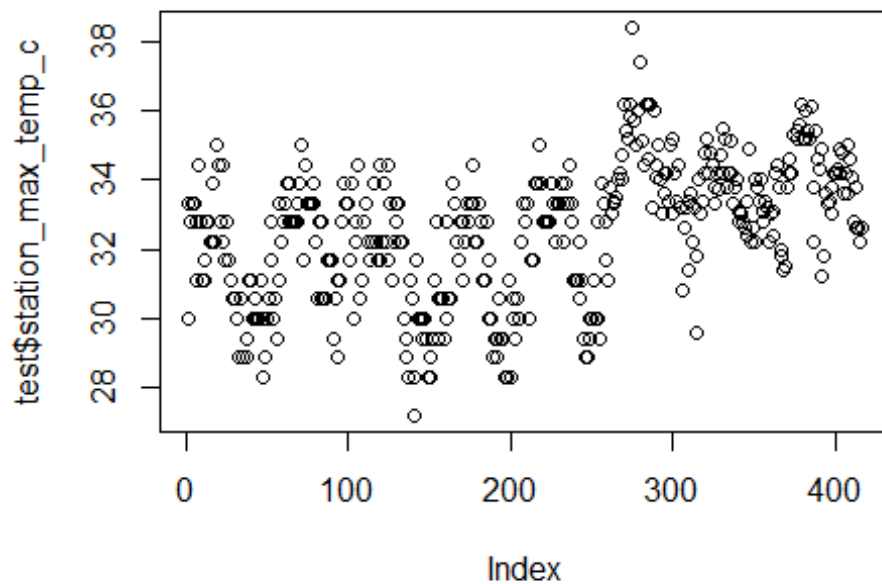


```
mean(test$station_diur_temp_rng_c, na.rm = TRUE)

## [1] 7.810991

test$station_diur_temp_rng_c[is.na(test$station_diur_temp_rng_c)] <-
mean(test$station_diur_temp_rng_c, na.rm = TRUE)

plot(test$station_max_temp_c)
```

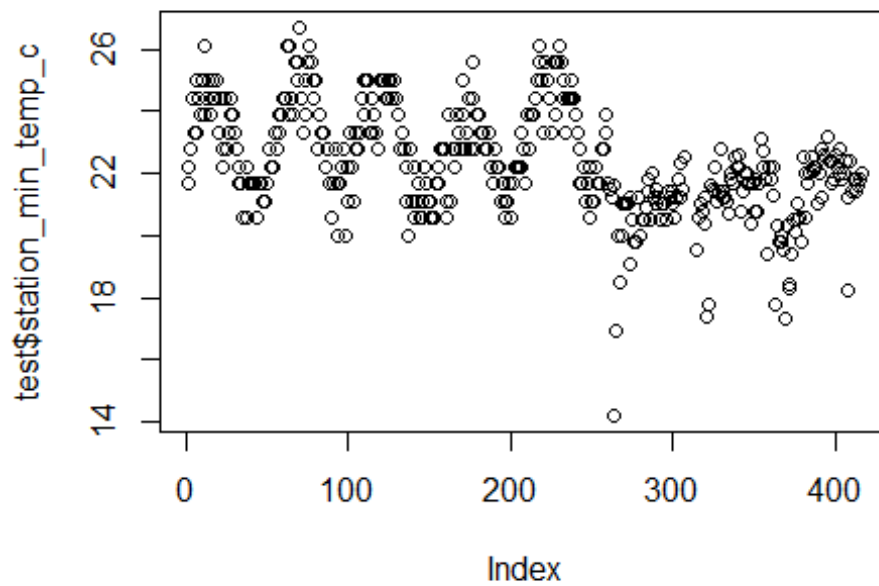


```
mean(test$station_max_temp_c, na.rm = TRUE)
```

```
## [1] 32.53462
```

```
test$station_max_temp_c[is.na(test$station_max_temp_c)] <-  
mean(test$station_max_temp_c, na.rm = TRUE)
```

```
plot(test$station_min_temp_c)
```

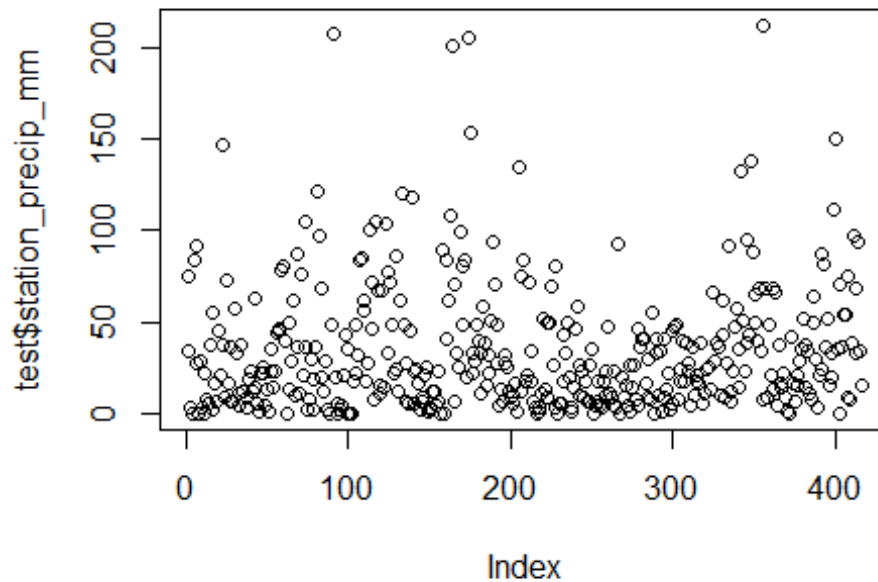


```
mean(test$station_min_temp_c, na.rm = TRUE)

## [1] 22.36855

test$station_min_temp_c[is.na(test$station_min_temp_c)] <-
mean(test$station_min_temp_c, na.rm = TRUE)

plot(test$station_precip_mm)
```



```
mean(test$station_precip_mm,na.rm = TRUE)

## [1] 34.27859

test$station_precip_mm[is.na(test$station_precip_mm)] <-
mean(test$station_precip_mm,na.rm = TRUE)

# count missing values (as percent)
apply(test, 2, function(x)
  round(100 * (length(which(is.na(x))))/length(x) , digits = 1)) %>%
  as.data.frame() %>%
  `names<-`('Percent of Missing Values')

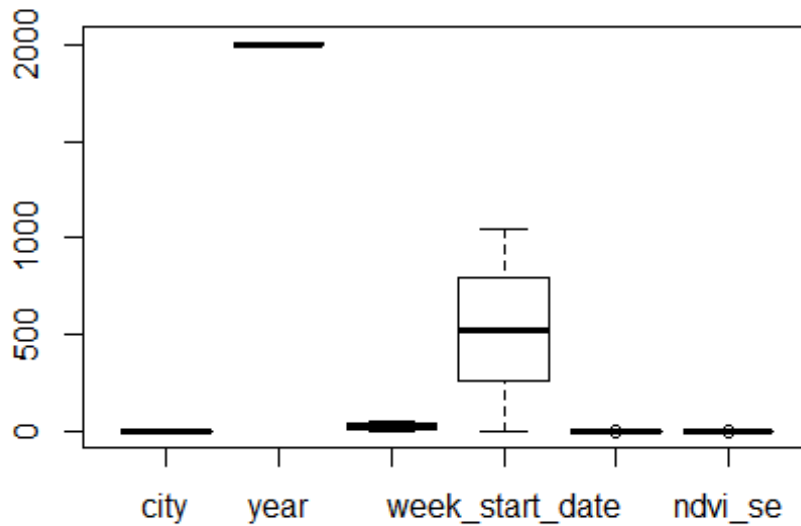
##                               Percent of Missing Values
## city                               0.0
## year                               0.0
## weekofyear                         0.0
## week_start_date                    0.0
## ndvi_nw                            0.0
## ndvi_se                            0.0
## ndvi_sw                            0.2
## precipitation_amt_mm               0.0
## reanalysis_air_temp_k              0.0
## reanalysis_avg_temp_k              0.0
## reanalysis_dew_point_temp_k        0.0
## reanalysis_max_air_temp_k          0.0
```

```

## reanalysis_min_air_temp_k 0.0
## reanalysis_precip_amt_kg_per_m2 0.0
## reanalysis_relative_humidity_percent 0.0
## reanalysis_sat_precip_amt_mm 0.0
## reanalysis_specific_humidity_g_per_kg 0.0
## reanalysis_tdtr_k 0.0
## station_avg_temp_c 0.0
## station_diur_temp_rng_c 0.0
## station_max_temp_c 0.0
## station_min_temp_c 0.0
## station_precip_mm 0.0

##Checking the Outliers for the Values
boxplot(train[1:6])

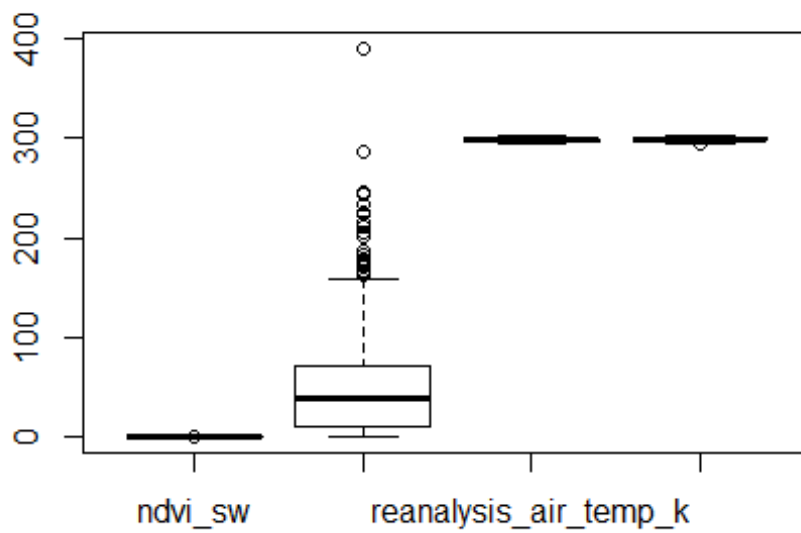
```



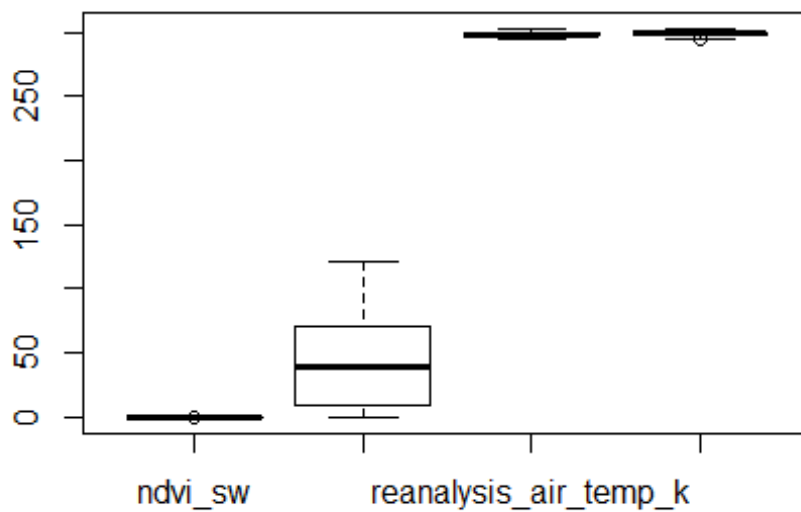
```

boxplot(train[7:10])

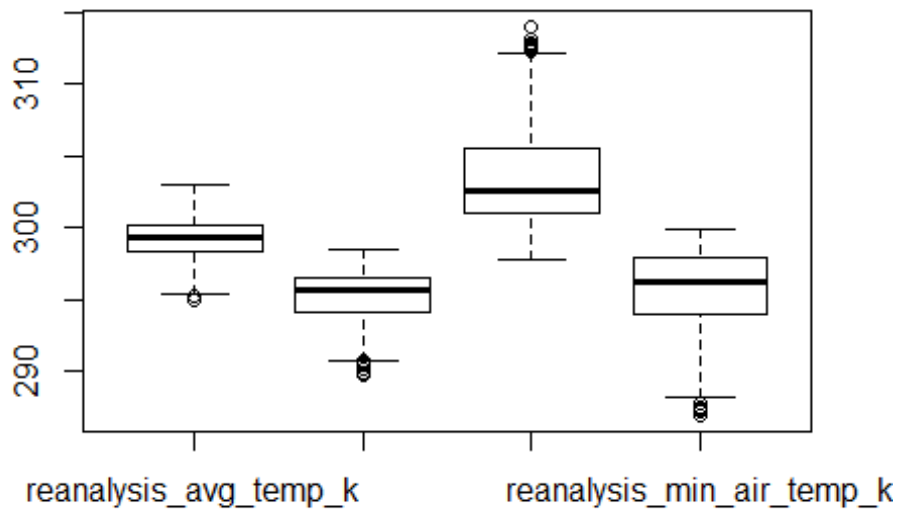
```



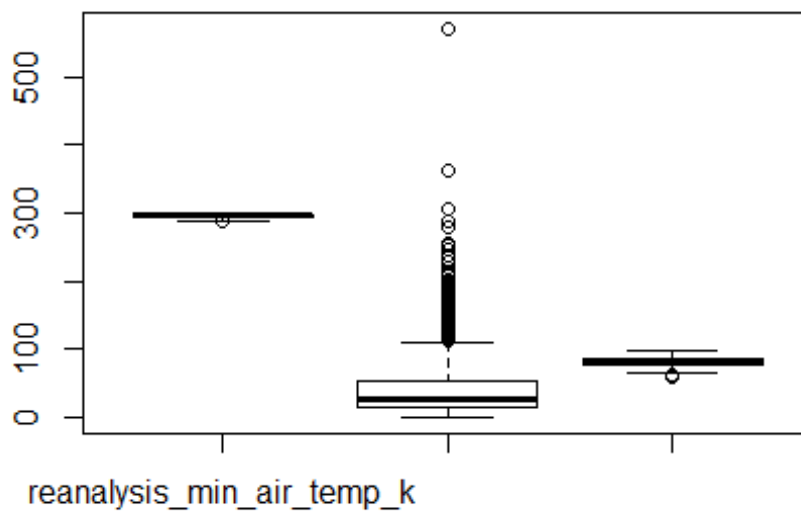
```
train$precipitation_amt_mm <- squish(train$precipitation_amt_mm,  
quantile(train$precipitation_amt_mm, c(.05, .95)))  
boxplot(train[7:10])
```




```
boxplot(train[10:13])
```



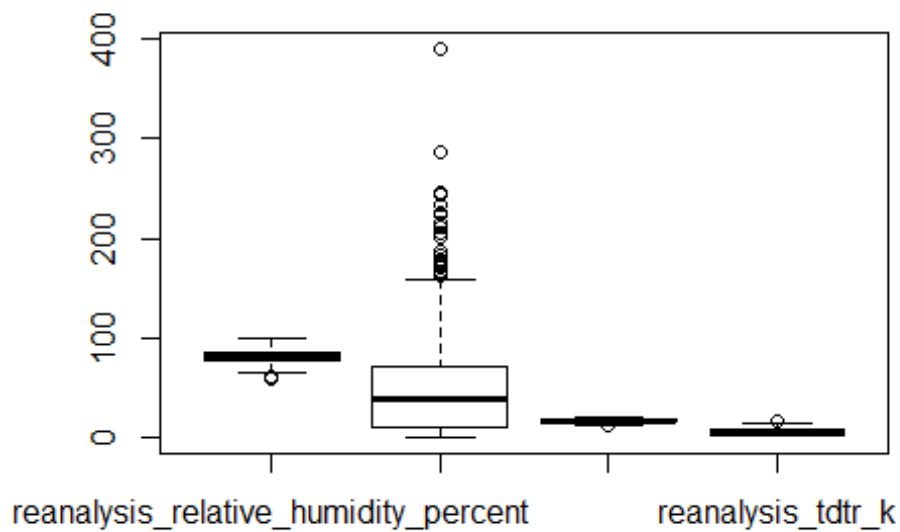
```
boxplot(train[13:15])
```



```

train$reanalysis_precip_amt_kg_per_m2 <-
squish(train$reanalysis_precip_amt_kg_per_m2,
quantile(train$reanalysis_precip_amt_kg_per_m2, c(.05, .95)))
boxplot(train[15:18])

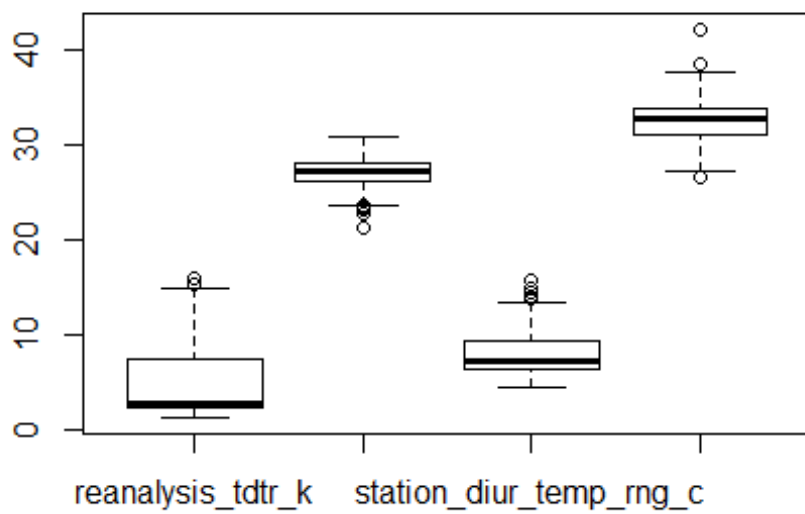
```



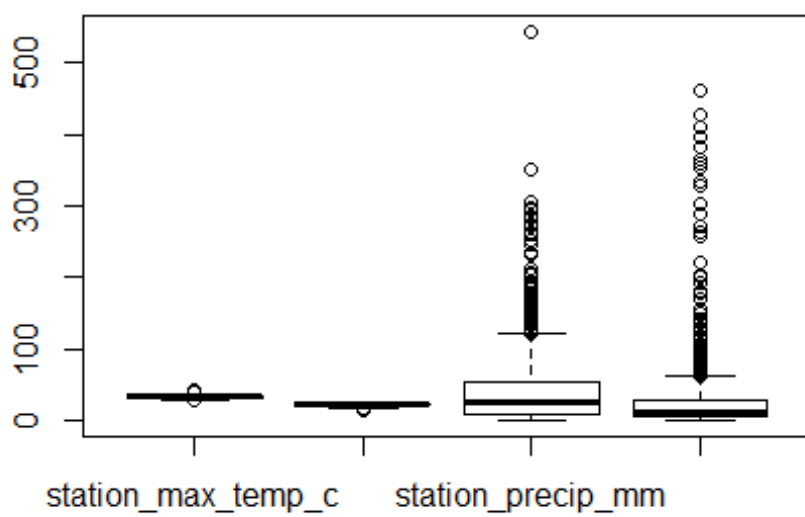
```

train$reanalysis_sat_precip_amt_mm <-
squish(train$reanalysis_sat_precip_amt_mm,
quantile(train$reanalysis_sat_precip_amt_mm, c(.05, .95)))
boxplot(train[18:21])

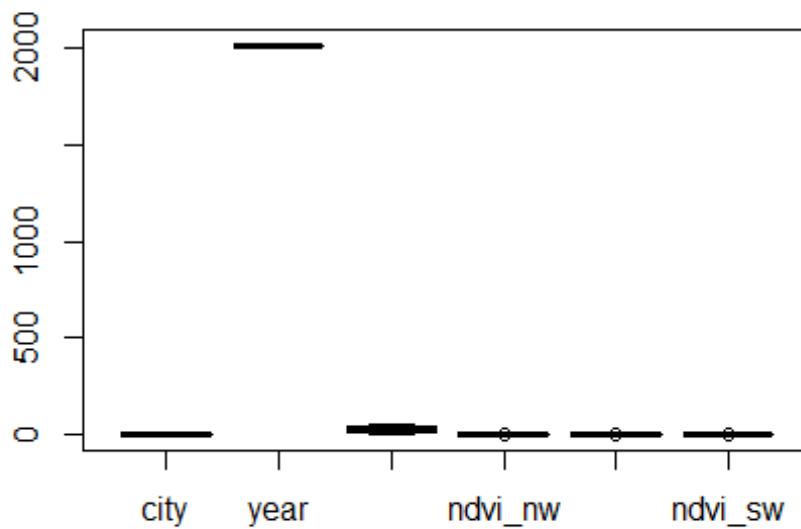
```



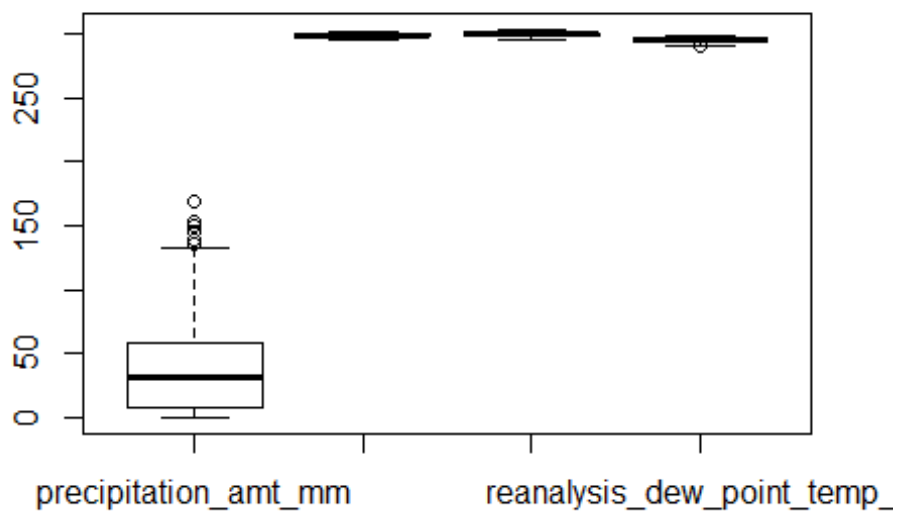
```
boxplot(train[21:24])
```



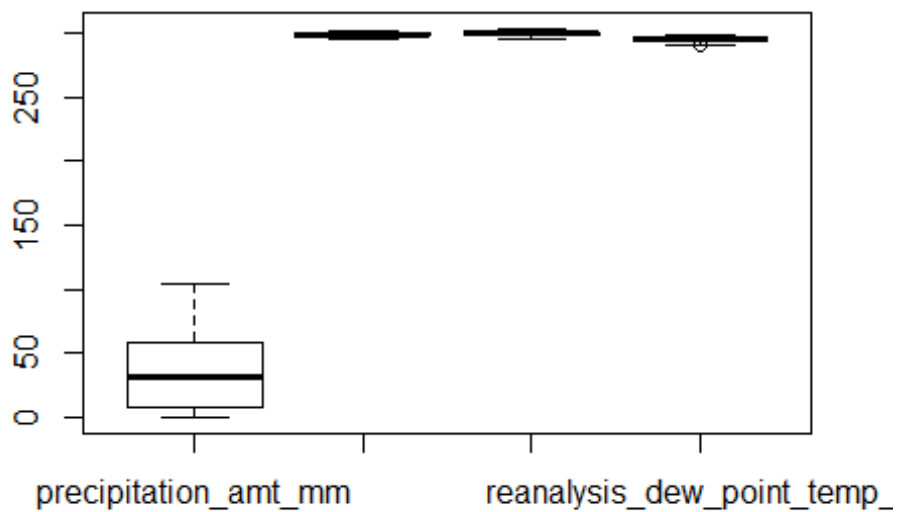
```
train$station_precip_mm <- squish(train$station_precip_mm,  
quantile(train$station_precip_mm, c(.05, .95)))  
  
train$week_start_date <- NULL  
test$week_start_date <- NULL  
  
##Checking the Outliers for the Values  
boxplot(test[1:6])
```



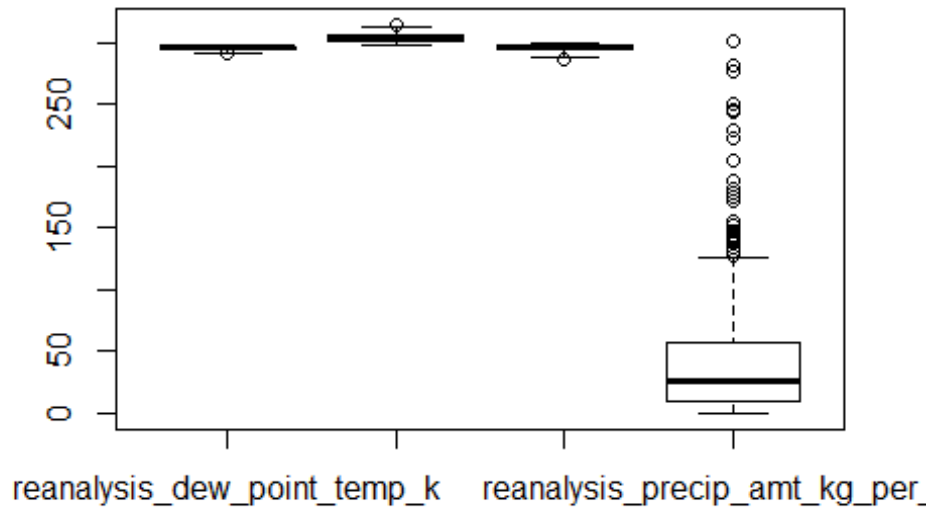
```
boxplot(test[7:10])
```



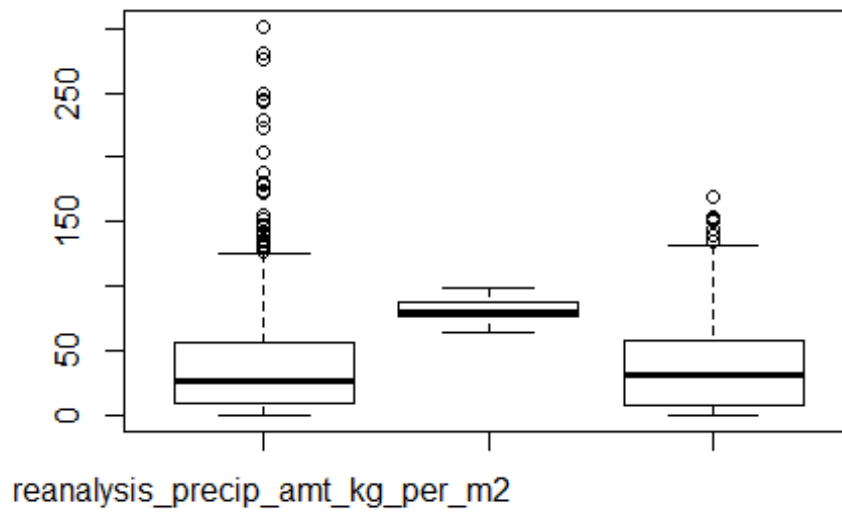
```
test$precipitation_amt_mm <- squish(test$precipitation_amt_mm,
quantile(test$precipitation_amt_mm, c(.05, .95)))
boxplot(test[7:10])
```



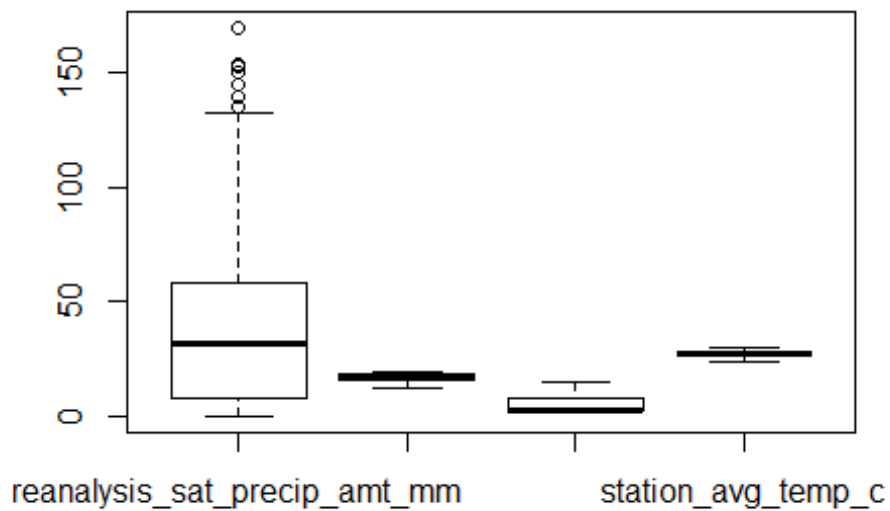
```
boxplot(test[10:13])
```



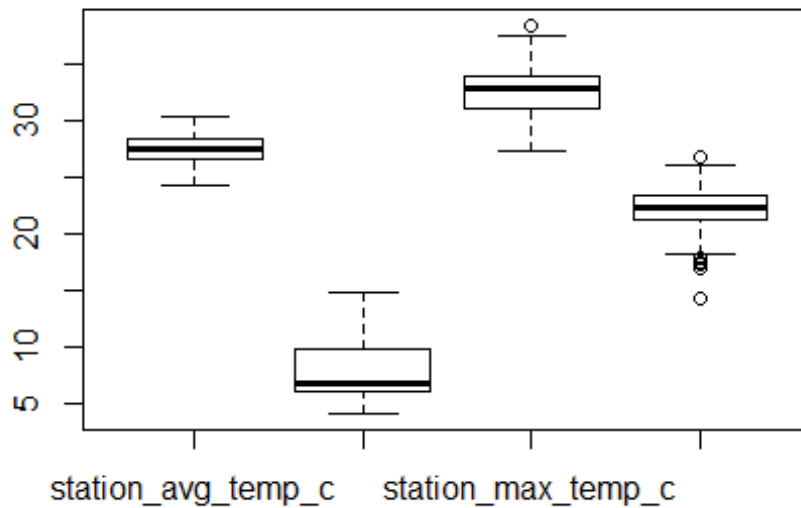
```
boxplot(test[13:15])
```



```
test$reanalysis_precip_amt_kg_per_m2 <-
squish(test$reanalysis_precip_amt_kg_per_m2,
quantile(test$reanalysis_precip_amt_kg_per_m2, c(.05, .95)))
boxplot(test[15:18])
```



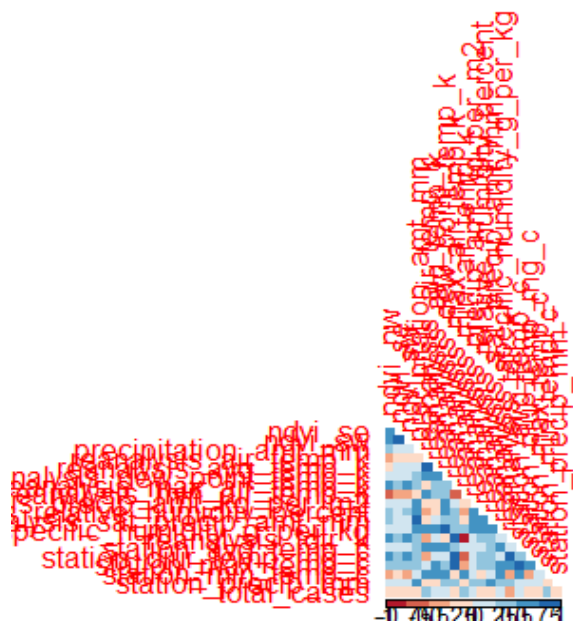
```
test$reanalysis_sat_precip_amt_mm <-
squish(test$reanalysis_sat_precip_amt_mm,
quantile(test$reanalysis_sat_precip_amt_mm, c(.05, .95)))
boxplot(test[18:21])
```



```
#boxplot(test[21:24])
test$station_precip_mm <- squish(test$station_precip_mm,
quantile(test$station_precip_mm, c(.05, .95)))

## Plotting Some Data and Getting the Result on the basis of the same
train %>%
  dplyr::select(-city, -year, -weekofyear) %>%
  cor(use = 'pairwise.complete.obs') -> M1

corrplot(M1, type="lower", method="color",
col=brewer.pal(n=8, name="RdBu"),diag=FALSE)
```

##Precisely none of these correlations are very strong.
 ##After seeing the Corelation we are moving forward to some of the Inputs

```
# reanalysis_specific_humidity_g_per_kg
#reanalysis_dew_point_temp_k
#station_avg_temp_c
#station_min_temp_c
#reanalysis_min_temp_k
#reanalysis_air_temp_k
```

```
#####
#                               Modelling                               #
#####
```

```
test_new  <- test[,c("reanalysis_air_temp_k" ,"reanalysis_min_air_temp_k",
"reanalysis_specific_humidity_g_per_kg", "reanalysis_dew_point_temp_k",
"station_avg_temp_c" , "station_min_temp_c")]
train_new <- train[,c("reanalysis_air_temp_k" ,"reanalysis_min_air_temp_k",
"reanalysis_specific_humidity_g_per_kg", "reanalysis_dew_point_temp_k",
"station_avg_temp_c" , "station_min_temp_c")]
train_pred <- train[,c("total_cases")]
```

```
fit <- glm.nb("total_cases ~ 1 + reanalysis_specific_humidity_g_per_kg +
reanalysis_air_temp_k + reanalysis_dew_point_temp_k + station_avg_temp_c +
station_min_temp_c" , data = train_new )
```

```

summary(train_new)

## reanalysis_air_temp_k reanalysis_min_air_temp_k
## Min. :294.6 Min. :286.9
## 1st Qu.:297.7 1st Qu.:293.9
## Median :298.7 Median :296.2
## Mean :298.7 Mean :295.7
## 3rd Qu.:299.8 3rd Qu.:297.9
## Max. :302.2 Max. :299.9
## reanalysis_specific_humidity_g_per_kg reanalysis_dew_point_temp_k
## Min. :11.72 Min. :289.6
## 1st Qu.:15.56 1st Qu.:294.1
## Median :17.07 Median :295.6
## Mean :16.75 Mean :295.2
## 3rd Qu.:17.97 3rd Qu.:296.5
## Max. :20.46 Max. :298.4
## station_avg_temp_c station_min_temp_c
## Min. :21.40 Min. :14.7
## 1st Qu.:26.33 1st Qu.:21.1
## Median :27.39 Median :22.2
## Mean :27.19 Mean :22.1
## 3rd Qu.:28.13 3rd Qu.:23.3
## Max. :30.80 Max. :25.6

sj_iq_test <- test[[1]]
sj_iq_test$predicted = predict(fit , test_new, type = 'response')

## Warning in sj_iq_test$predicted = predict(fit, test_new, type =
## "response"): Coercing LHS to a list

fin_pred_val <- round(sj_iq_test$predicted)

submissions = read.csv("D:\\Kaggle Projects\\Dengue\\submission_format.csv",
header=TRUE, sep=",")

submissions$total_cases <- fin_pred_val

write.csv(submissions, 'D:\\Kaggle Projects\\Dengue\\predictions.csv',
row.names = FALSE)

```