mestrado-renata-ml

Carregando os pacotes exigidos

```
library(readx1)
library(tidyverse)
library(geobr)
library(skimr)
library(tidymodels)
library(ISLR)
library(modeldata)
library(wip)
library(ggpubr)
source("R/my_functions.R")
```

Listando os arquivos com os mapas de cada área separadamente

```
files_eu <- list.files("data/EU espacial/",full.names = TRUE)
files_sp <- list.files("data/SP espacial/",full.names = TRUE)</pre>
```

Carregando os mapa para Eucalipto

```
eu <- map_df(files_eu,grd_read)
```

Arquivo com os dados de emissão, temperatura e umidade (temporal)

Arquivo com os dados dos atributos do solo, somente

geoespacializados

Unindo as bases de dados, ou seja, repetindo os dados do solo para cada dia de avaliação

```
data_eu <- left_join(temporal_eu, spatial_eu, by="id") %>%
  select(-numero) %>%
  mutate(data = make_date(year= ano, month=mes, day=dia)) %>%
  relocate(id,data)
```

Carregando os mapa para Sistema Silvipastoril

```
sp <- map_df(files_sp,grd_read)</pre>
```

Arquivo com os dados de emissão, temperatura e umidade (temporal)

Arquivo com os dados dos atributos do solo, somente geoespacializados

Unindo as bases de dados, ou seja, repetindo os dados do solo

para cada dia de avaliação

```
data_sp <- left_join(temporal_sp, spatial_sp, by="id") %>%
  select(-numero) %>%
  mutate(data = make_date(year= ano, month=mes, day=dia)) %>%
  relocate(id,data)
```

```
tibble(xnames=names(data_eu), ynames=names(data_sp)) %>%
 mutate(logic test = xnames == ynames)
#> # A tibble: 21 × 3
     xnames ynames logic_test
   <chr> <chr> <lgl>
#>
#> 1 id id
                  TRUE
#> 2 data data TRUE
#> 3 ano ano TRUE
\#> 4 mes mes TRUE
#> 5 dia dia TRUE
\#> 6 F F TRUE \#> 7 T TRUE
U #> 9 Al A<sup>1</sup> #> 10
                  TRUE
          AI
                   TRUE
                   TRUE
#> # i 11 more rows
```

Unindo toda a base de dados

Aprendizado de Máquina

```
# Definindo a base de treino e a base de teste
locais <- data set %>% pull(local) %>% unique()
for(i in seq along(locais)){
 lo <- locais[i]</pre>
 data <- data set %>% filter(local == lo)
 dias <- data$data %>% unique()
 for(j in seq along(dias)){
   di <- dias[j]</pre>
    df <- data %>% filter(data == di)
    fco2 initial split <- initial split(df %>%
select(-c(id, ano, mes, dia, local)) %>%
                                                                                  sample n
(trunc(nrow(df)*.51289)), prop = 0.75)
    fco2_train <- training(fco2_initial_split)</pre>
    hist_fco2 <- fco2_train %>%
      ggplot(aes(x=F, y=..density..))+
      geom histogram(bins = 30, color="black", fill="lightgray") +
      geom density(alpha=.05, fill="red") +
      theme bw() +
      labs(x="FCO2", y = "Densidade", title = paste(lo, di))
    print(hist fco2)
    fco2 train %>%
      select(where(is.numeric)) %>%
      drop na() %>%
      cor() %>%
      corrplot::corrplot()
    fco2 recipe <- recipe(F ~ ., data = fco2 train) %>%
      step_novel(all_nominal_predictors()) %>%
      step zv(all predictors()) %>%
      step dummy(all nominal predictors())
    bake(prep(fco2 recipe), new data = NULL)
    # visdat::vis miss(bake(prep(fco2 recipe), new data = NULL))
    fco2 resamples <- vfold cv(fco2 train, v = 5) #<-----
    ### DECISION TREE
    print("ARVORE DE DECISÃO")
    fco2 dt model <- decision tree(</pre>
     cost_complexity = tune(),
     tree depth = tune(),
     min n = tune()
    ) 응>응
      set mode("regression") %>%
      set engine("rpart")
    fco2 dt wf <- workflow()</pre>
      add model(fco2 dt model) %>%
```

```
add recipe(fco2 recipe)
 grid dt <- grid random(</pre>
   cost complexity(c(-6, -4)),
   tree depth(range = c(8, 18)),
   min n(range = c(42, 52)),
   size = 2 # <-----
 fco2_dt_tune_grid <- tune_grid(</pre>
   fco2 dt wf,
   resamples = fco2 resamples,
   grid = grid dt,
   metrics = metric set(rmse)
 print(autoplot(fco2 dt tune grid))
 fco2 dt best params <- select best(fco2 dt tune grid, "rmse")
 fco2 dt wf <- fco2 dt wf %>% finalize workflow(fco2 dt best params)
 fco2 dt last fit <- last fit(fco2 dt wf, fco2 initial split)
 fco2 test preds <- bind rows(
   collect predictions(fco2 dt last fit) %>% mutate(modelo = "dt")
 pre obs plot <- fco2 test preds %>%
   ggplot(aes(x=.pred, y=F)) +
   geom point()+
   theme bw() +
   geom smooth(method = "lm") +
   stat regline equation(ggplot2::aes(
     label = paste(..eq.label.., ..rr.label.., sep = "*plain(\",\")~~")))+
   labs(title = paste(lo,di))
 print(pre obs plot)
 fco2 modelo final <- fco2 dt wf %>% fit(df)
 saveRDS(fco2 modelo final,
          paste0("models-3/fco2 modelo dt ",lo," ",di,".rds"))
 fco2 dt last fit model <- fco2 dt last fit$.workflow[[1]]$fit$fit</pre>
 vip plot <- vip(fco2 dt last fit model,</pre>
      aesthetics = list(color = "grey35", size = 0.8, fill="orange")) +
   theme_bw()
 print(vip plot)
 da <- fco2 test preds %>%
filter(F > 0, .pred>0)
 my r <- cor(da$F,da$.pred)</pre>
 my r2 <- my r*my r
 my mse <- Metrics::mse(da$F,da$.pred)</pre>
 my rmse <- Metrics::rmse(da$F,</pre>
 my mae <- Metrics::mae(da$F,da$.pred)</pre>
```

```
my mape <- Metrics::mape(da$F,da$.pred)*100
vector of metrics <- c(r=my r, R2=my r2, MSE=my mse,
                       RMSE=my rmse, MAE=my mae, MAPE=my mape)
print(data.frame(vector of metrics))
# ##RANDOM FOREST
print("RANDOM FOREST")
fco2 rf model <- rand forest(</pre>
 min_n = tune(),
 mtry = tune(),
 trees = tune()
  응>응
  set mode("regression") %>%
  set engine("randomForest")
fco2 rf wf <- workflow()</pre>
  add model(fco2 rf model) %>%
  add recipe(fco2 recipe)
grid rf <- grid random(</pre>
 min_n(range = c(20, 30)),
 mtry(range = c(5, 10)),
 trees(range = c(100, 500)),
  size = 2
fco2 rf tune grid <- tune grid(
 fco2 rf wf,
 resamples = fco2 resamples,
 grid = grid rf,
 metrics = metric set(rmse)
print(autoplot(fco2 rf tune grid))
fco2 rf best params <- select best(fco2 rf tune grid, "rmse")
fco2 rf wf <- fco2 rf wf %>% finalize workflow(fco2 rf best params)
fco2 rf last fit <- last fit(fco2 rf wf, fco2 initial split)</pre>
fco2 test preds <- bind rows(</pre>
  collect predictions(fco2 rf last fit) %>% mutate(modelo = "rf")
pre_obs_plot <- fco2_test_preds %>%
 ggplot(aes(x=.pred, y=F)) +
  geom point()+
 theme bw() +
 geom smooth(method = "lm") +
  stat regline equation(ggplot2::aes(
   label = paste(..eq.label.., ..rr.label.., sep = "*plain(\",\")~~"))+
  labs(title = paste(lo,di))
print(pre obs plot)
fco2 modelo final <- fco2 rf wf %>% fit(df)
```

```
saveRDS(fco2 modelo final,
          paste0("models-3/fco2 modelo rf ",lo," ",di,".rds"))
 fco2 rf last fit model <- fco2 rf last fit$.workflow[[1]]$fit$fit</pre>
 vip plot <- vip(fco2 rf last fit model,</pre>
      aesthetics = list(color = "grey35", size = 0.8, fill="orange")) +
    theme bw()
 print(vip_plot)
      da <- fco2 test preds %>%
filter(F > 0, .pred>0)
 my r <- cor(da$F,da$.pred)</pre>
 my_r2 \leftarrow my_r*my_r
 my mse <- Metrics::mse(da$F,da$.pred)</pre>
 my rmse <- Metrics::rmse(da$F,</pre>
                            da$.pred)
 my mae <- Metrics::mae(da$F,da$.pred)</pre>
 my mape <- Metrics::mape(da$F,da$.pred)*100</pre>
 vector_of_metrics <- c(r=my_r, R2=my_r2, MSE=my_mse,</pre>
                          RMSE=my rmse, MAE=my mae, MAPE=my mape)
 print(data.frame(vector_of_metrics))
  ##XGBOOST
  cores = 6
  fco2 xgb_model <- boost_tree(</pre>
   mtry = 0.8,
    trees = tune(), # <-----</pre>
   min n = 5,
    tree depth = 4,
    loss_reduction = 0, # lambda
    learn rate = tune(), # epsilon
    sample size = 0.8
  ) %>%
    set mode("regression") %>%
    set engine("xgboost", nthread = cores, counts = FALSE)
 fco2 xgb wf <- workflow()</pre>
    add model(fco2 xgb model) %>%
    add recipe(fco2 recipe)
 grid xgb <- expand.grid(</pre>
    learn_rate = c(0.05, 0.3),
    trees = c(2, 250, 500)
  )
  #passo 1
  fco2 xgb tune grid <- tune grid(
    fco2 xgb wf,
    resamples = fco2_resamples,
    grid = grid xgb,
    metrics = metric set(rmse)
```

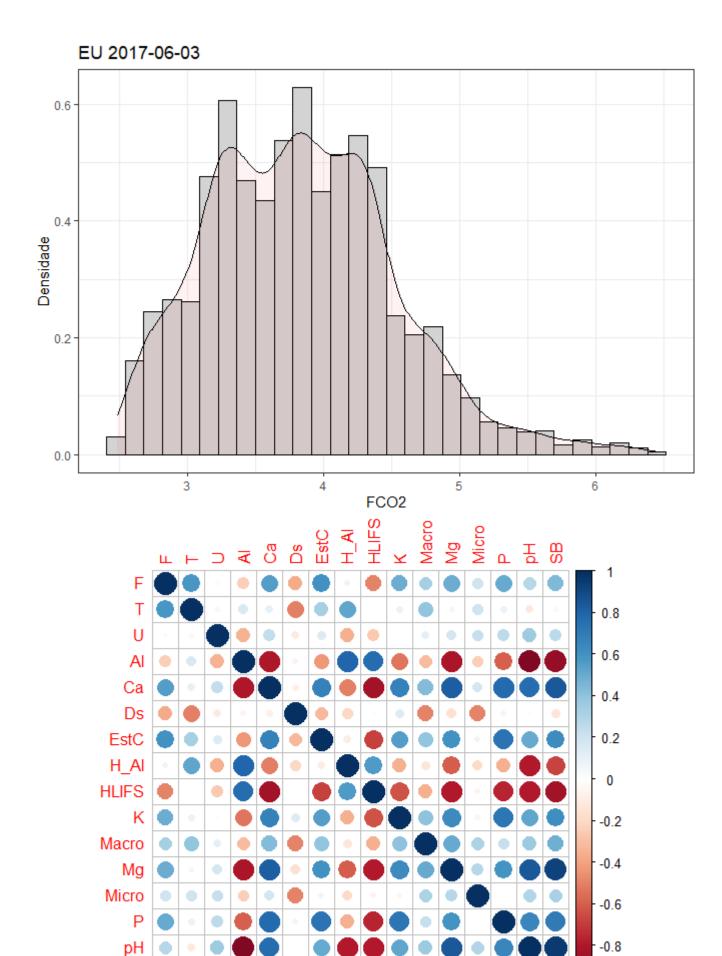
```
# print(autoplot(fco2 xgb tune grid))
    fco2 xgb select best passo1 <- fco2 xgb tune grid %>%
      select_best(metric = "rmse")
    # passo 2
   fco2 xgb model <- boost tree(</pre>
     mtry = 0.8,
      trees = fco2_xgb_select_best_passo1$trees,
     min n = tune(),
      tree_depth = tune(),
      loss reduction = 0,
     learn rate = fco2 xgb select best passo1$learn rate,
      sample size = 0.8
   ) 응>응
      set mode("regression") %>%
      set engine("xgboost", nthread = cores, counts = FALSE)
   fco2 xgb wf <- workflow() %>%
      add model(fco2 xgb model) %>%
      add recipe(fco2 recipe)
   fco2_xgb_grid <- expand.grid(</pre>
     tree depth = c(1, 3, 4),
     min n = c(5, 30, 60)
    fco2 xgb tune grid <- fco2 xgb wf %>%
      tune grid(
       resamples =fco2 resamples,
        grid = fco2 xgb grid,
        control = control grid(save pred = TRUE, verbose = FALSE, allow par = TRUE),
       metrics = metric set(rmse)
    fco2 xgb select best passo2 <- fco2 xgb tune grid %>% select best(metric = "rms
e")
    # passo 3
    fco2 xgb model <- boost tree(</pre>
     mtry = 0.8,
     trees = fco2 xgb select best passo1$trees,
     min n = fco2 xgb select best passo2$min n,
      tree depth = fco2 xgb select best passo2$tree depth,
      loss reduction =tune(),
      learn rate = fco2 xgb select best passo1$learn rate,
      sample size = 0.8
   ) 응>응
      set mode("regression") %>%
      set engine("xgboost", nthread = cores, counts = FALSE)
   fco2 xgb wf <- workflow() %>%
      add model(fco2 xgb model) %>%
      add recipe(fco2 recipe)
```

```
fco2 xgb grid <- expand.grid(</pre>
      loss reduction = c(0.01, 0.05, 1, 2, 4, 8)
    fco2_xgb_tune_grid <- fco2_xgb_wf %>%
     tune grid(
        resamples = fco2_resamples,
        grid = fco2 xgb grid,
        control = control_grid(save_pred = TRUE, verbose = FALSE, allow par = TRUE),
       metrics = metric set(rmse)
    fco2 xgb select best passo3 <- fco2 xgb tune grid %>% select best(metric = "rmse")
    # passo 4
    fco2 xgb model <- boost tree(</pre>
     mtry = tune(),
     trees = fco2 xgb select best passo1$trees,
     min_n = fco2_xgb_select_best_passo2$min_n,
     tree depth = fco2 xgb select best passo2$tree depth,
      loss reduction = fco2 xgb select best passo3$loss reduction,
      learn rate = fco2 xgb select best passo1$learn rate,
      sample size = tune()
   ) 응>응
      set mode("regression") |>
      set_engine("xgboost", nthread = cores, counts = FALSE)
      fco2 xgb wf <- workflow() %>%
      add model(fco2 xgb model) %>%
      add recipe(fco2 recipe)
   fco2 xgb grid <- expand.grid(</pre>
      sample size = seq(0.5, 1.0, length.out = 2), ## <---
     mtry = seq(0.1, 1.0, length.out = 2) ## <---
   )
    fco2 xgb tune grid <- fco2 xgb wf
     tune grid(
       resamples = fco2 resamples,
       grid = fco2 xgb grid,
        control = control grid(save pred = TRUE, verbose = FALSE, allow par = TRUE),
       metrics = metric_set(rmse)
    fco2_xgb_select_best_passo4 <- fco2_xgb_tune_grid %>% select best(metric = "rms
e")
    # passo 5
    fco2 xgb model <- boost tree(</pre>
     mtry = fco2_xgb_select_best_passo4$mtry,
     trees = tune(),
     min n = fco2 xgb select best passo2$min n,
      tree depth = fco2 xgb select best passo2$tree depth,
```

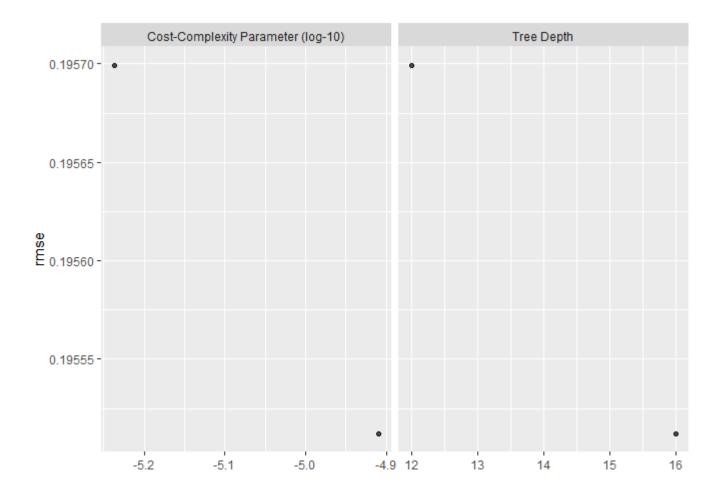
Grid

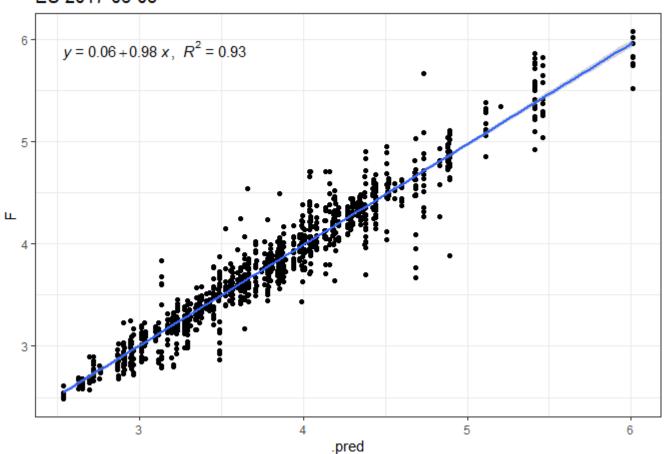
```
loss reduction = fco2 xgb select best passo3$loss reduction,
      learn rate = tune(),
      sample size = fco2 xgb select best passo4$sample size
    set_mode("regression") %>%
      set engine("xgboost", nthread = cores, counts = FALSE)
   fco2 xgb wf <- workflow() %>%
      add_model(fco2_xgb_model) %>%
      add recipe(fco2 recipe)
   fco2 xgb grid <- expand.grid(</pre>
     learn rate = c(0.05, 0.10, 0.15, 0.25),
      trees = c(100, 250, 500)
   )
    fco2 xgb tune grid <- fco2 xgb wf
      tune grid(
       resamples = fco2 resamples,
        grid = fco2 xgb grid,
        control = control grid(save pred = TRUE, verbose = FALSE, allow par = TRUE),
       metrics = metric set(rmse)
    fco2 xgb select best passo5 <- fco2 xgb tune grid %>% select best(metric = "rms
e")
    ## modelos final desempenho
    fco2 xgb model <- boost tree(</pre>
     mtry = fco2 xgb select best passo4$mtry,
     trees = fco2 xgb select best passo5$trees,
     min n = fco2 xgb select best passo2$min n,
      tree depth = fco2 xgb select best passo2$tree depth,
      loss reduction = fco2 xgb select best passo3$loss reduction,
      learn rate = fco2 xgb select best passo5$learn rate,
      sample_size = fco2_xgb_select_best_passo4$sample size
    ) 응>응
      set mode("regression") %>%
      set engine("xgboost", nthread = cores, counts = FALSE)
    df par <- data.frame(</pre>
      mtry = fco2 xgb select best passo4$mtry,
      trees = fco2 xgb select best passo5$trees,
     min n = fco2 xgb select best passo2$min n,
      tree depth = fco2 xgb select best passo2$tree depth,
      loss reduction = fco2 xgb select best passo3$loss reduction,
      learn rate = fco2 xgb select best passo5$learn rate,
      sample_size = fco2_xgb_select_best_passo4$sample_size
    fco2 xgb wf <- fco2 xgb wf %>% finalize workflow(df par) # <-----
    fco2 xgb last fit <- last fit(fco2 xgb wf, fco2 initial split)
```

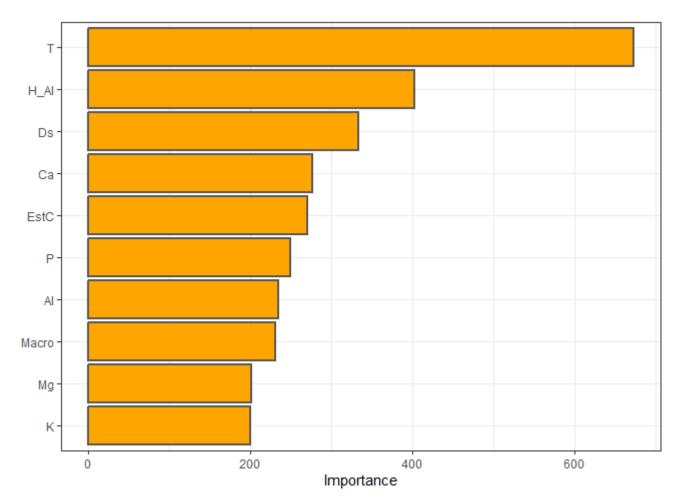
```
fco2 test preds <- bind rows(</pre>
      collect predictions(fco2 xgb last fit) %>% mutate(modelo = "xgb")
    pre obs plot <- fco2 test preds %>%
      ggplot(aes(x=.pred, y=F)) +
      geom point()+
      theme bw() +
      geom smooth(method = "lm") +
      stat_regline_equation(ggplot2::aes(
        label = paste(..eq.label.., ..rr.label.., sep = "*plain(\",\")~~")))+
      labs(title = paste(lo,di))
    print(pre obs plot)
    fco2 modelo final <- fco2 xgb wf %>% fit(df)
    saveRDS(fco2 modelo final,
            paste0("models-3/fco2_modelo_xgb_",lo,"_",di,".rds"))
    fco2_xgb_last_fit_model <- fco2_xgb_last_fit$.workflow[[1]]$fit$fit</pre>
    vip plot <- vip(fco2 xgb last fit model,</pre>
        aesthetics = list(color = "grey35", size = 0.8, fill="orange")) +
      theme bw()
    print(vip_plot)
        da <- fco2 test preds %>%
  filter(F > 0, .pred>0)
    my r <- cor(da$F,da$.pred)</pre>
    my r2 <- my r*my r
    my mse <- Metrics::mse(da$F,da$.pred)</pre>
    my rmse <- Metrics::rmse(da$F,</pre>
                              da$.pred)
    my_mae <- Metrics::mae(da$F,da$.pred)</pre>
    my_mape <- Metrics::mape(da$F,da$.pred)*100</pre>
    vector_of_metrics <- c(r=my_r, R2=my_r2, MSE=my_mse,</pre>
                            RMSE=my rmse, MAE=my mae, MAPE=my mape)
    print(data.frame(vector of metrics))
  }
}
```

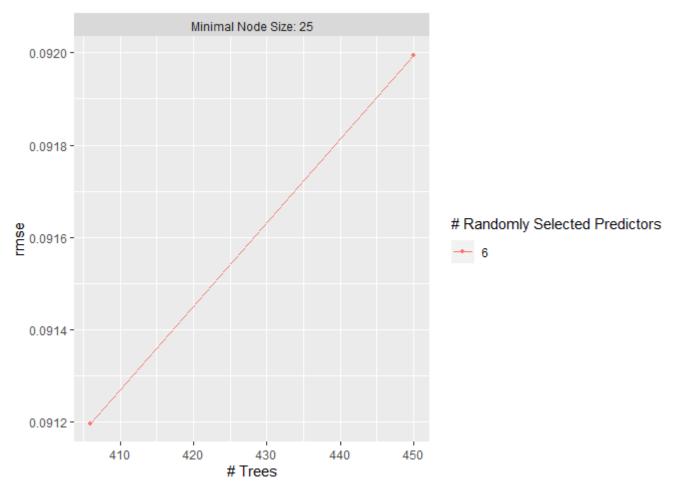


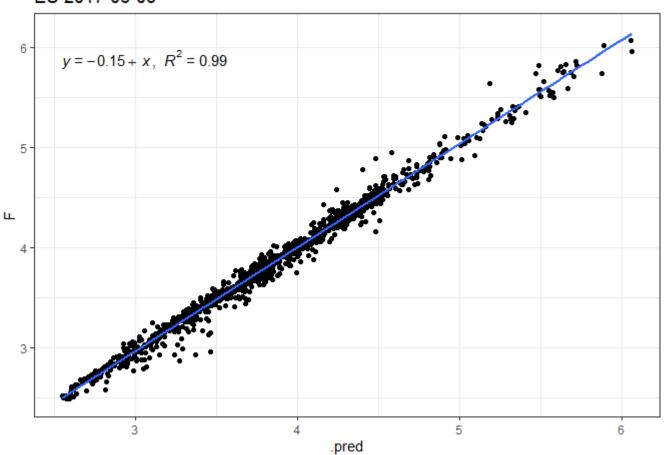
SB

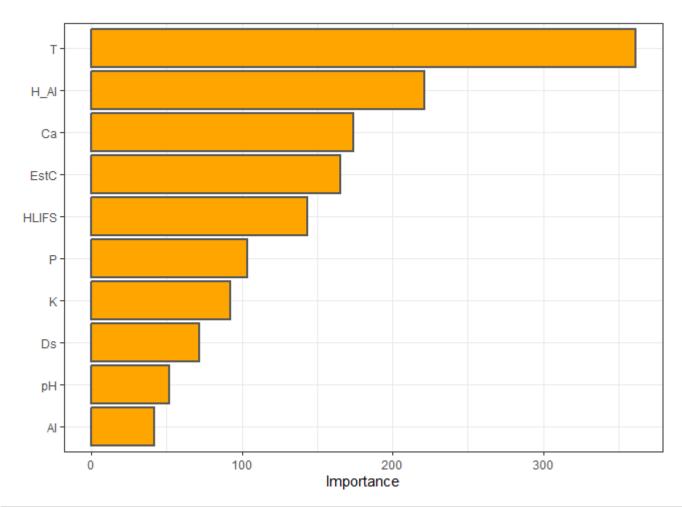


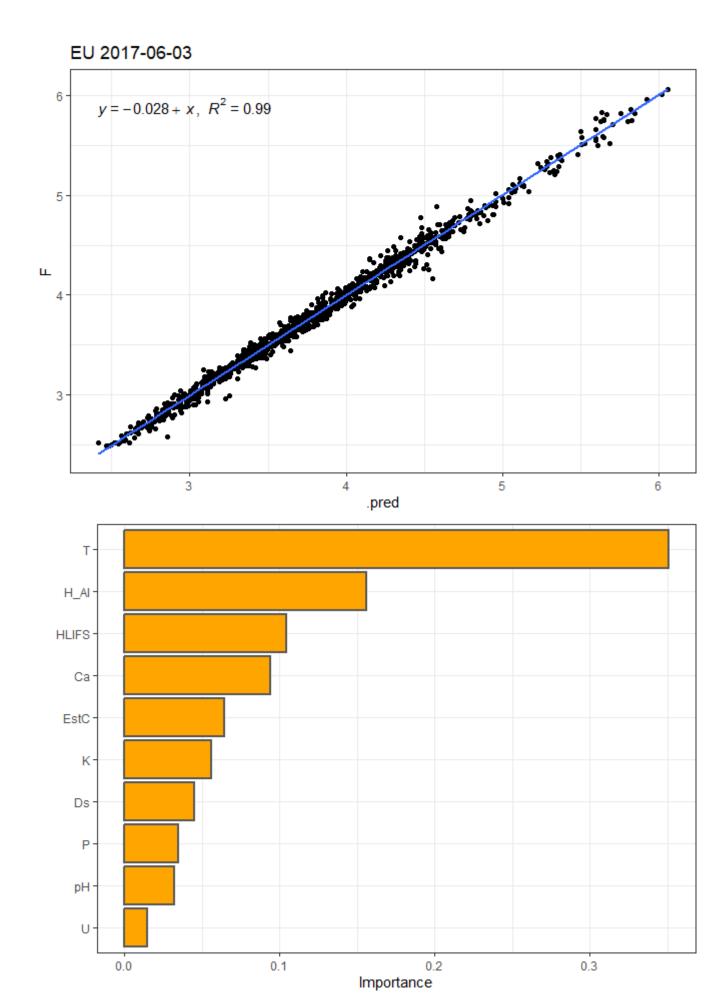


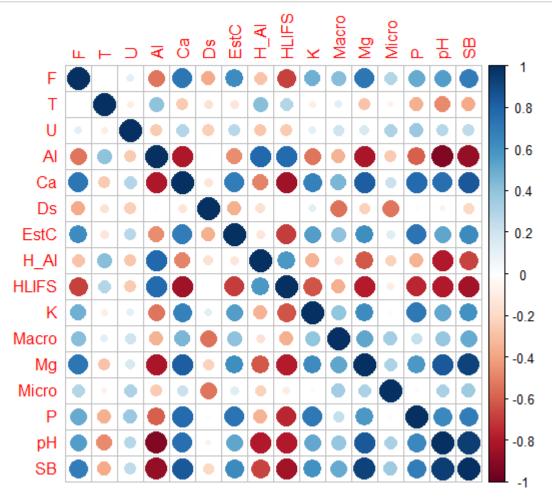




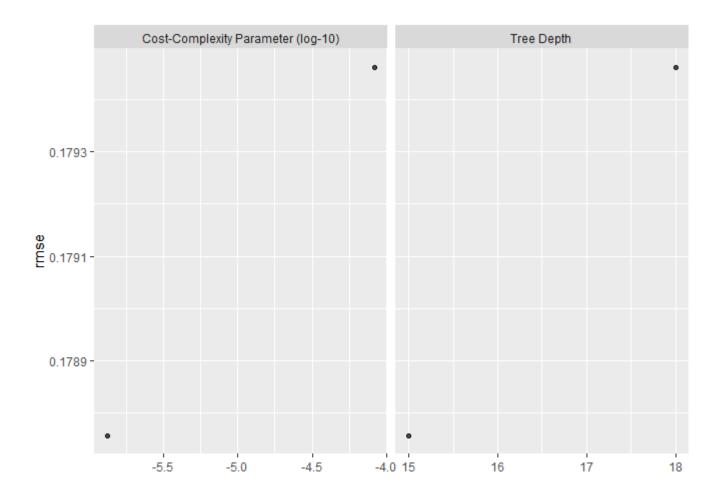


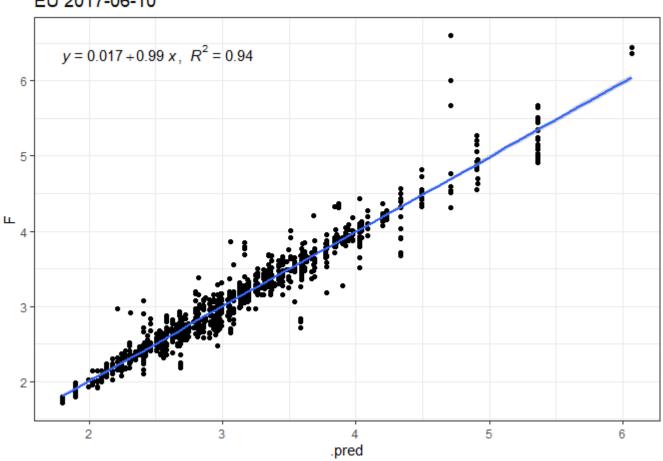


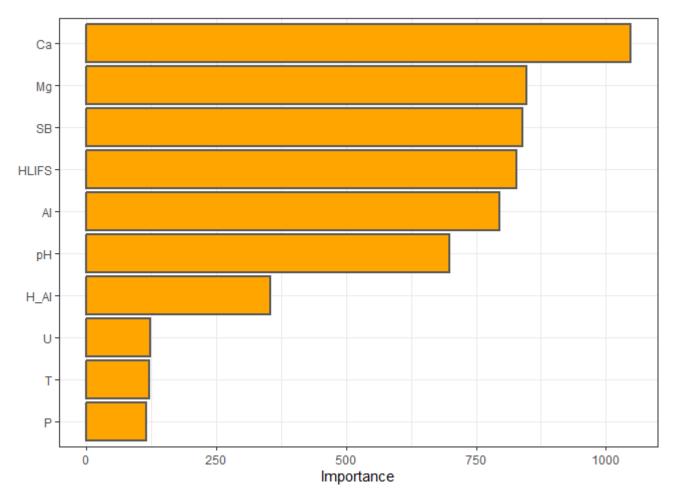


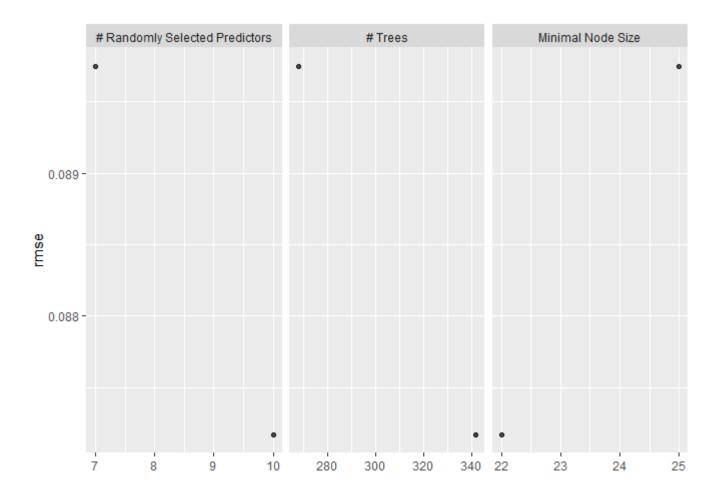


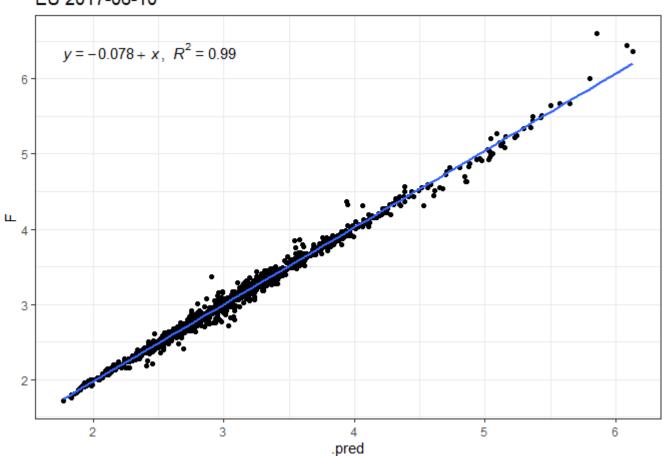
#> [1] "ARVORE DE DECISÃO"

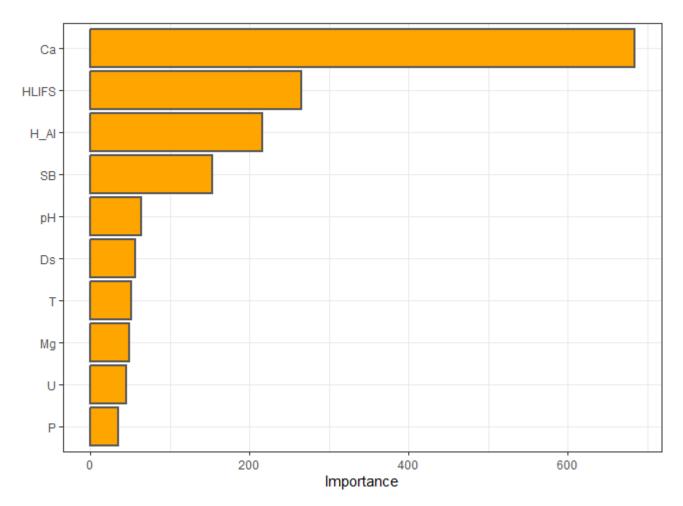




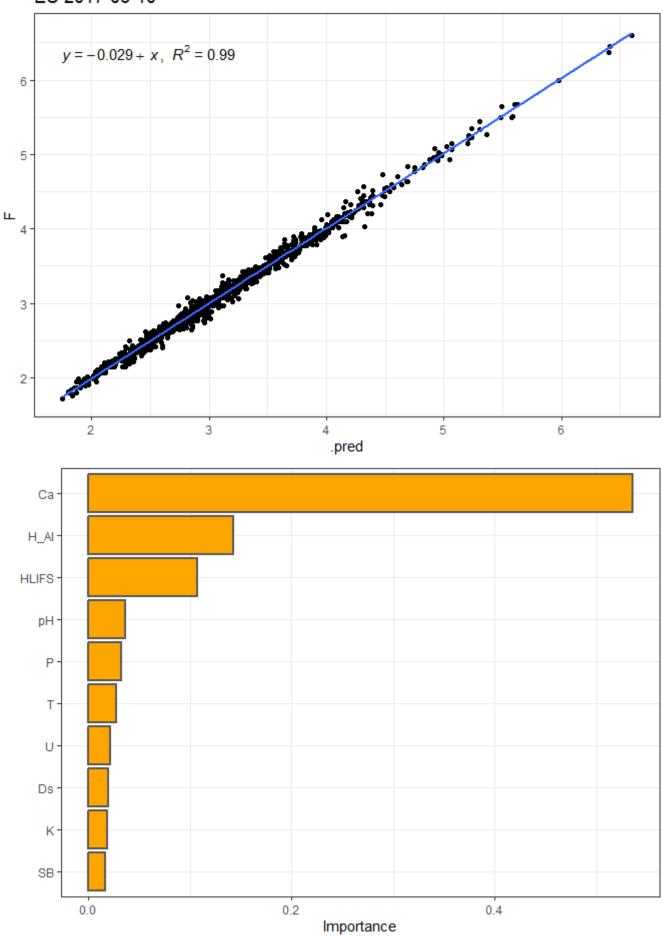


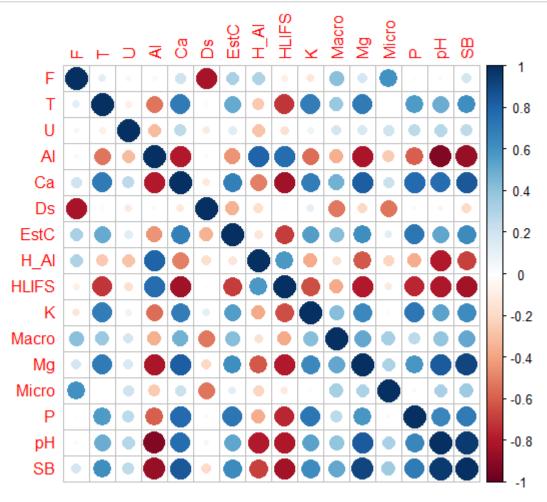




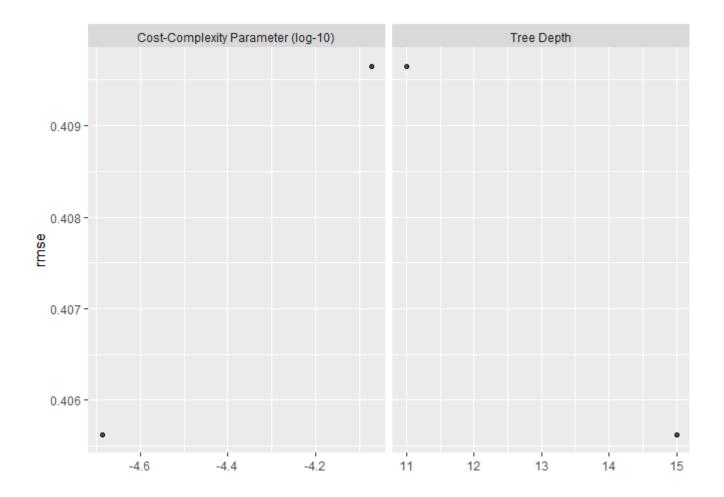


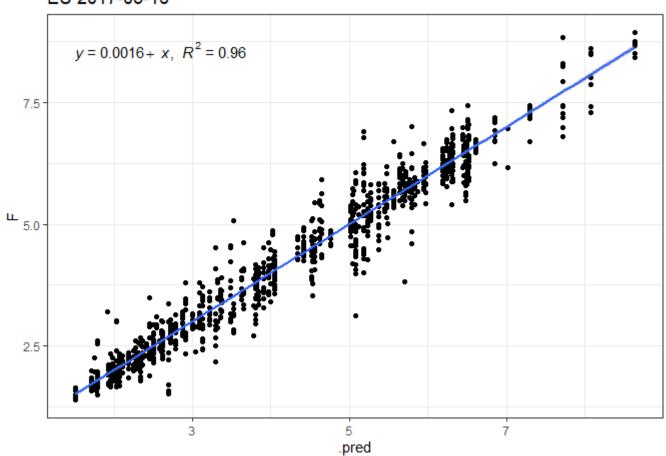


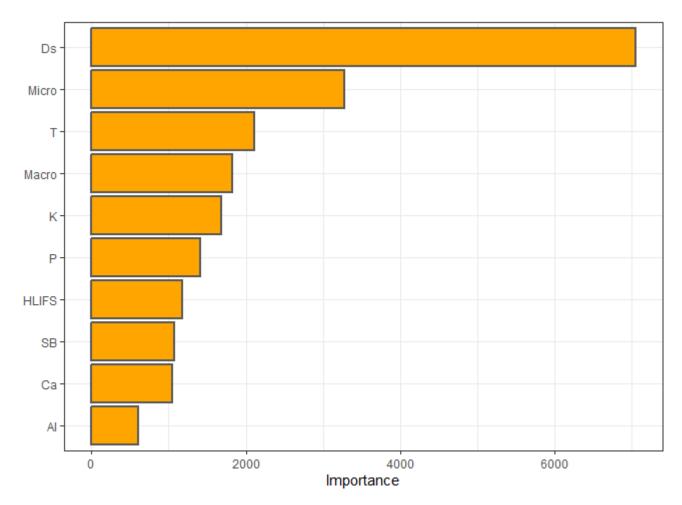


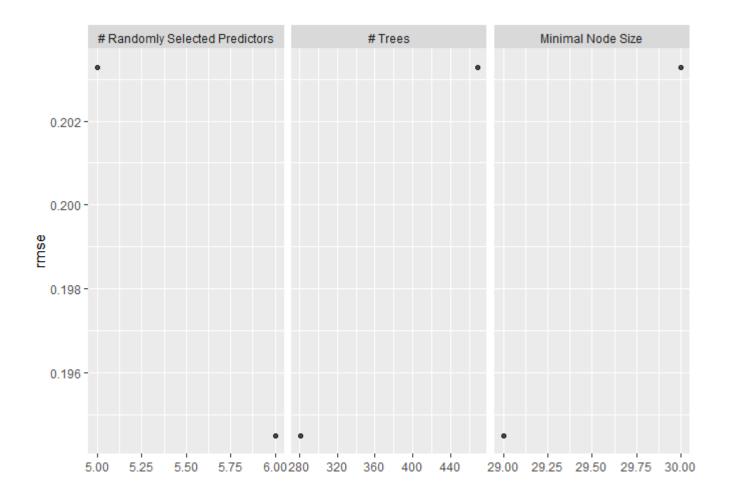


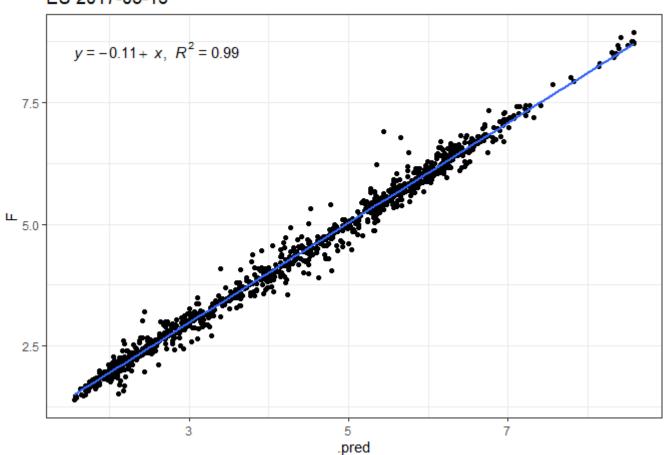
#> [1] "ARVORE DE DECISÃO"

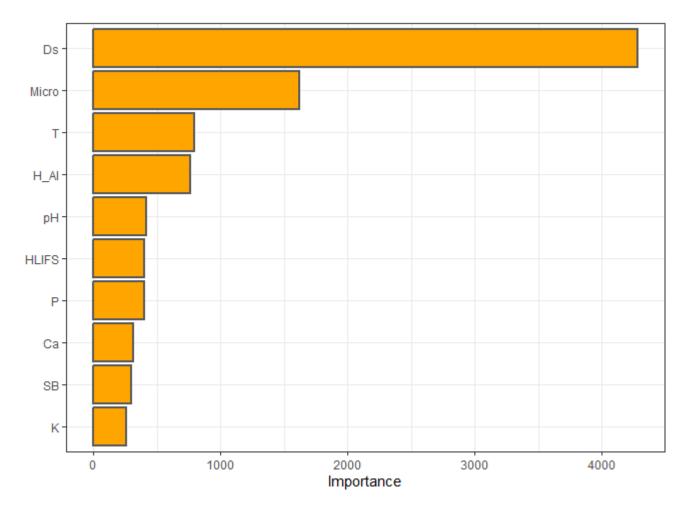




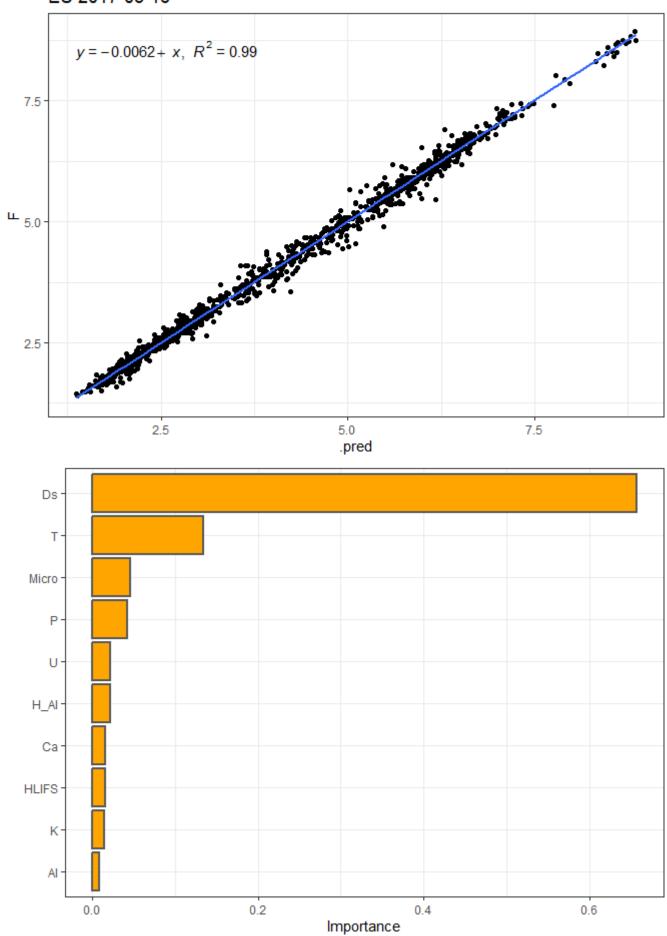




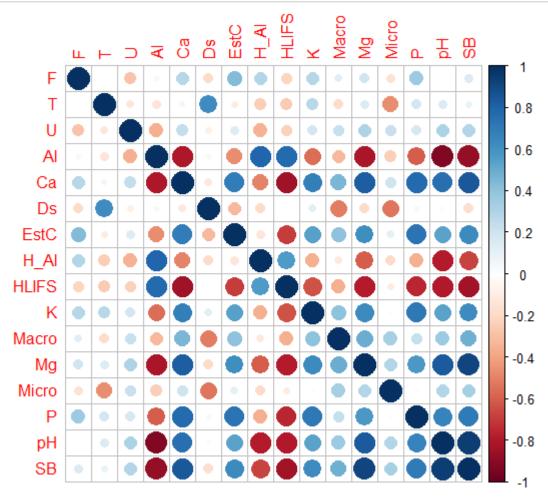




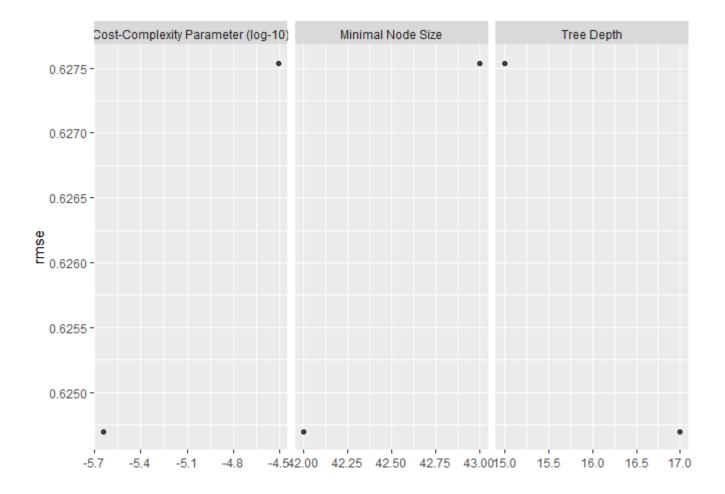


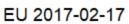


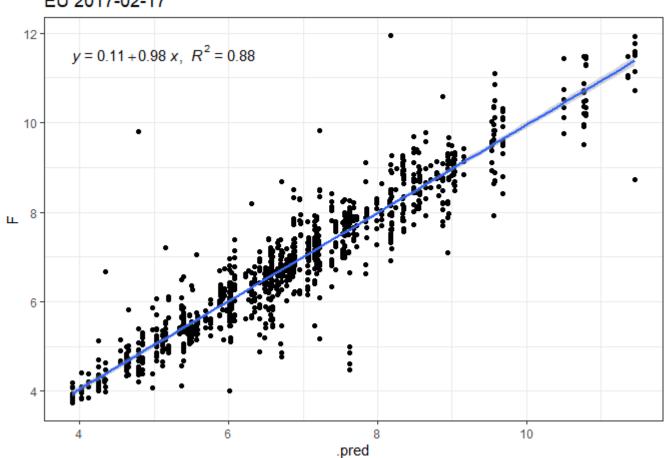
| #> | vector_of_metri |
|---------|-----------------|
| #> r | 0.99680631 |
| #> R2 | 0.99362282 |
| #> MSE | 0.02025893 |
| #> RMSE | 0.14233389 |
| #> MAE | 0.10093043 |
| #> MAPE | 2.69793849 |

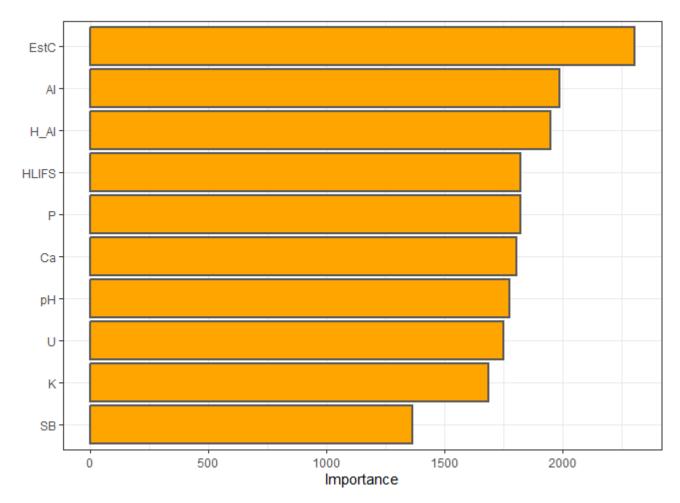


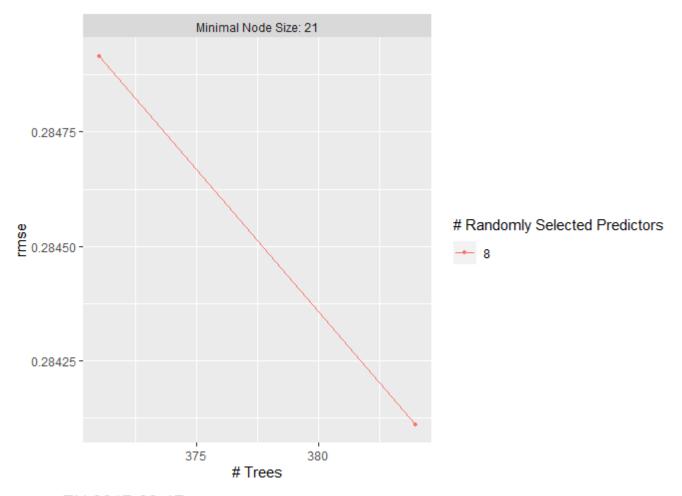
#> [1] "ARVORE DE DECISÃO"

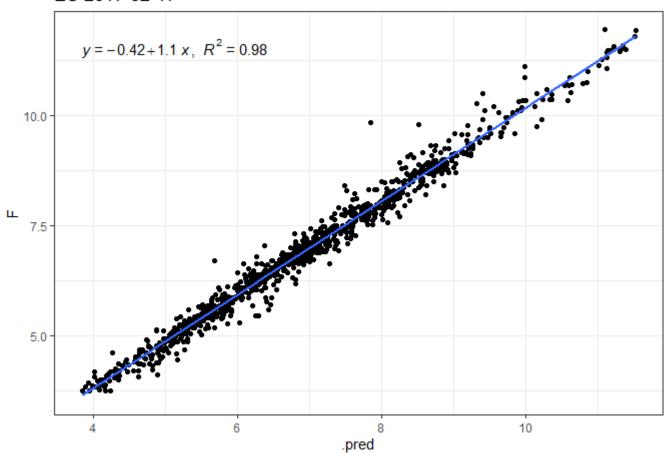


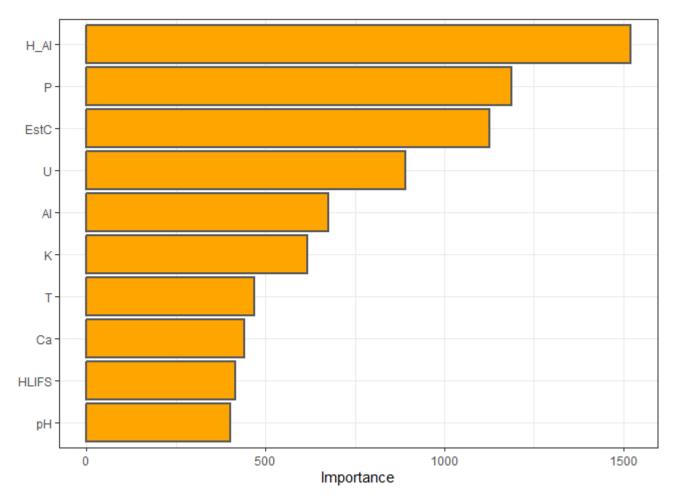




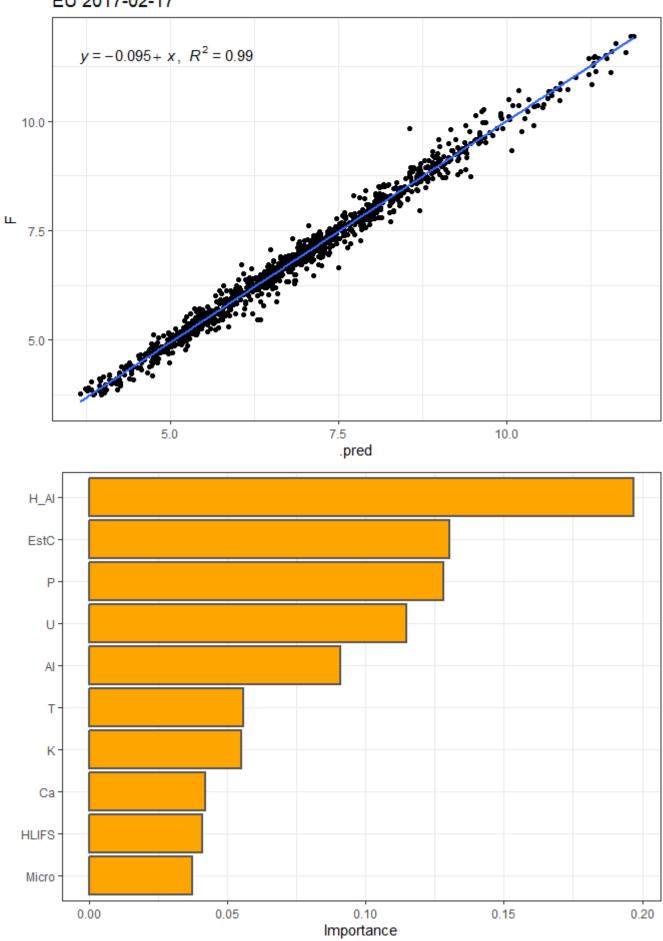




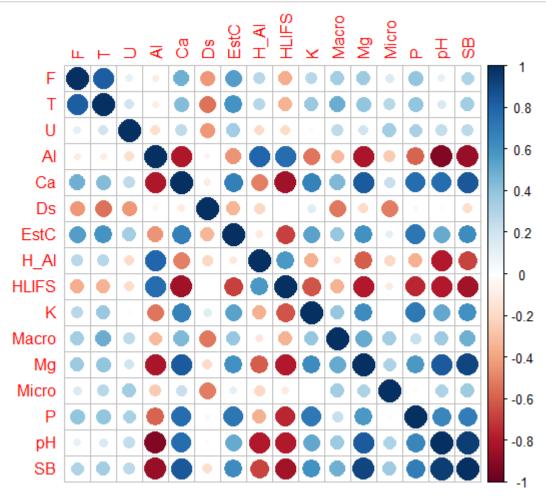


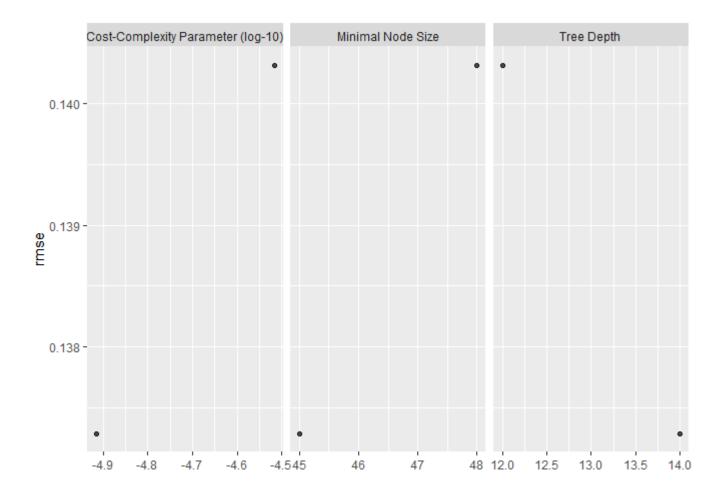




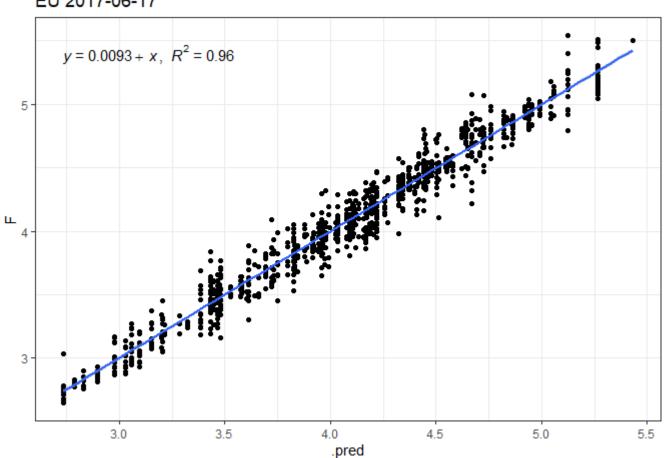


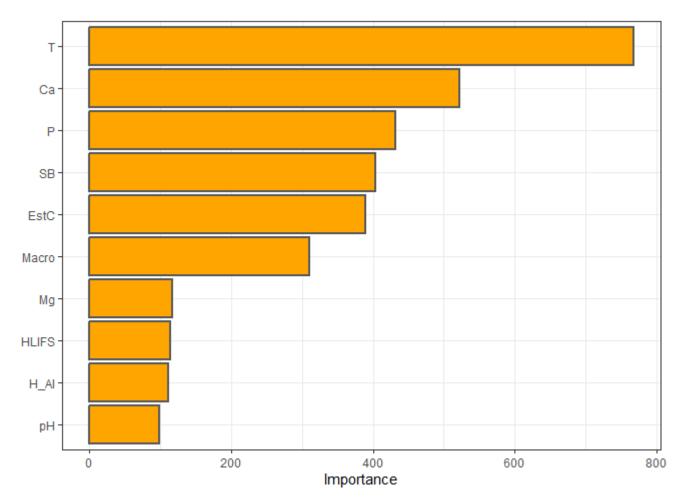
| #> | vector_of_metrics |
|---------|-------------------|
| #> r | 0.99263061 |
| #> R2 | 0.98531552 |
| #> MSE | 0.03929455 |
| #> RMSE | 0.19822853 |
| #> MAE | 0.14270433 |
| #> MAPE | 2.16523663 |
| | |

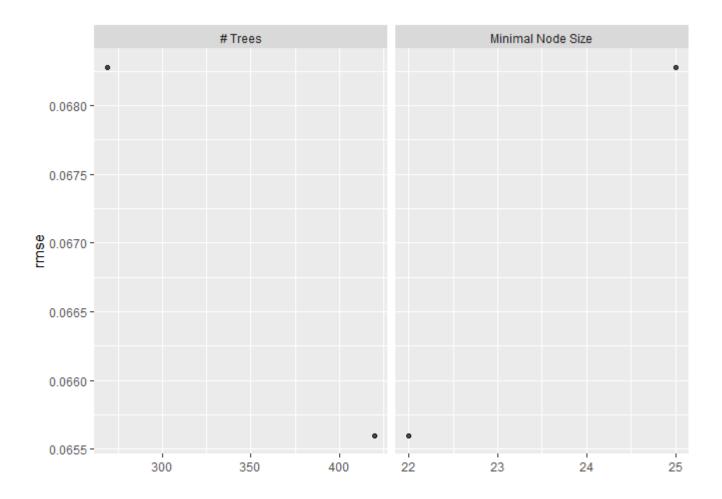




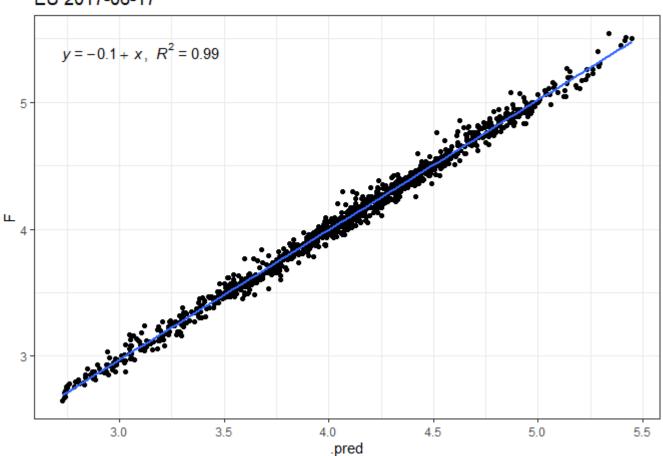
EU 2017-06-17

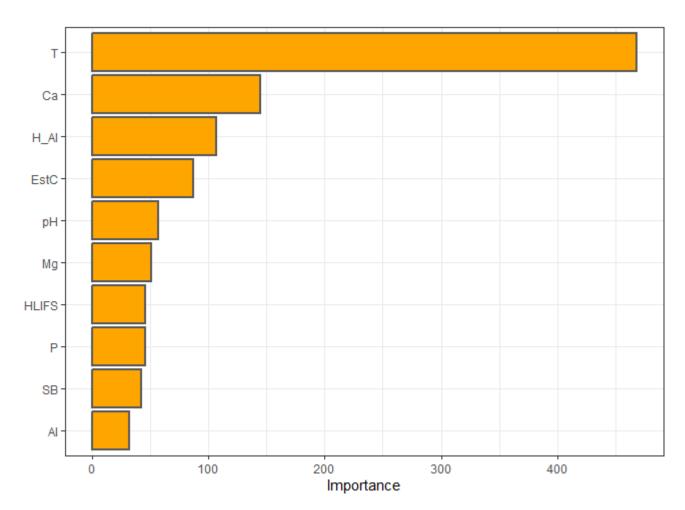




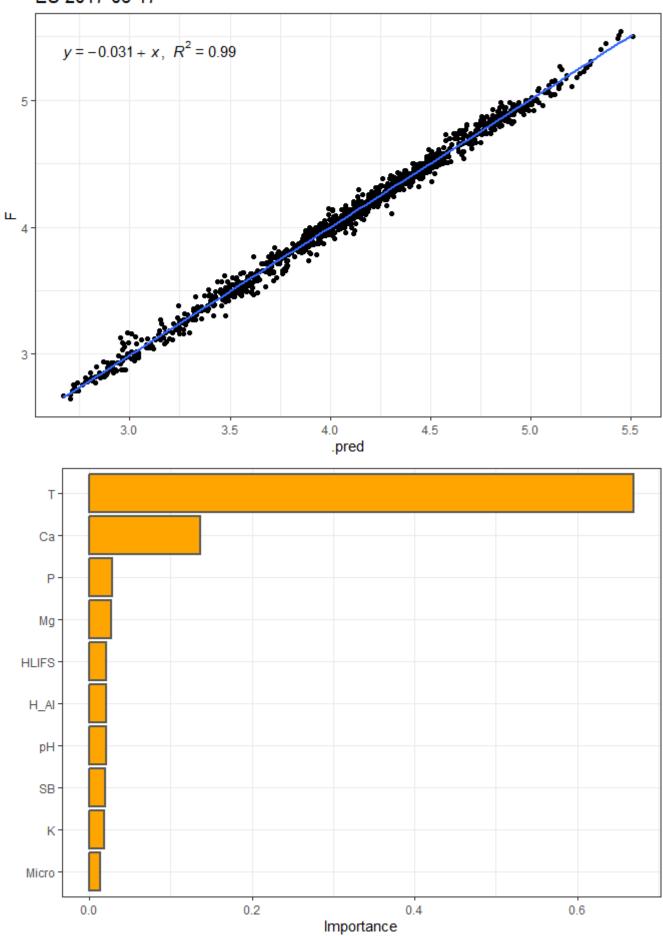


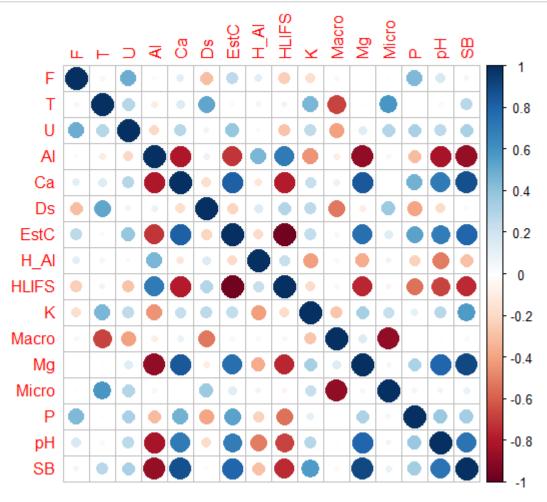
EU 2017-06-17

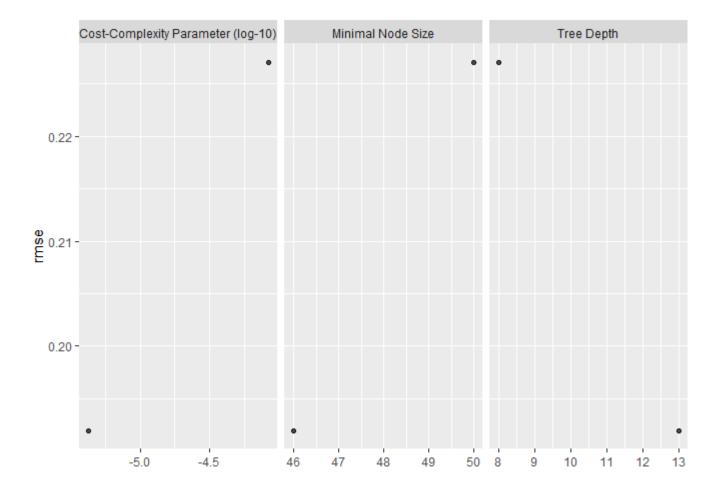


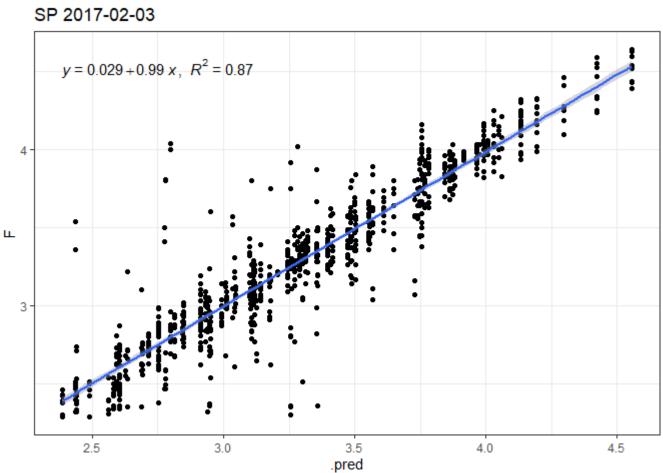


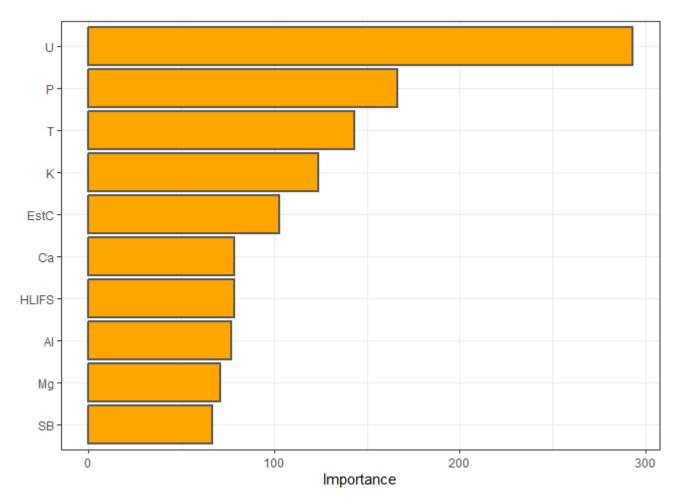


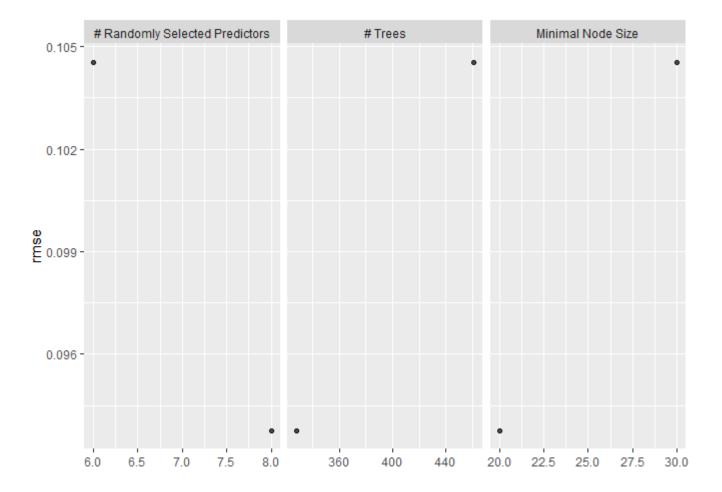


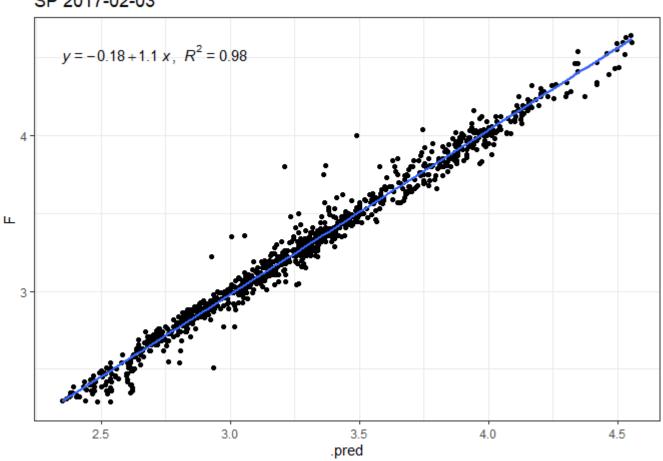


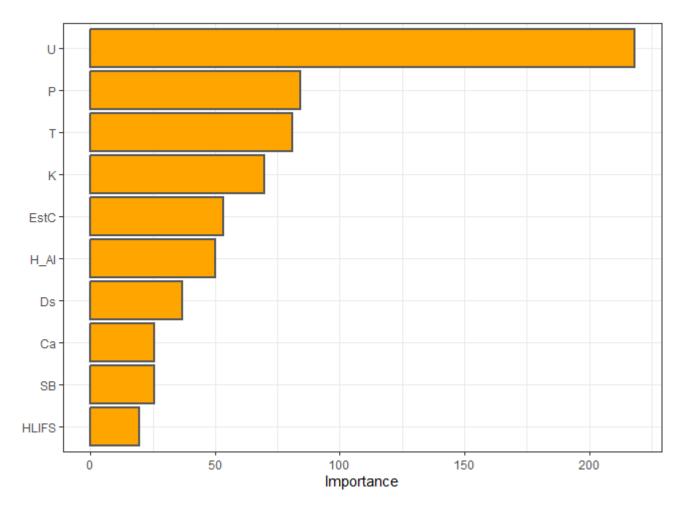


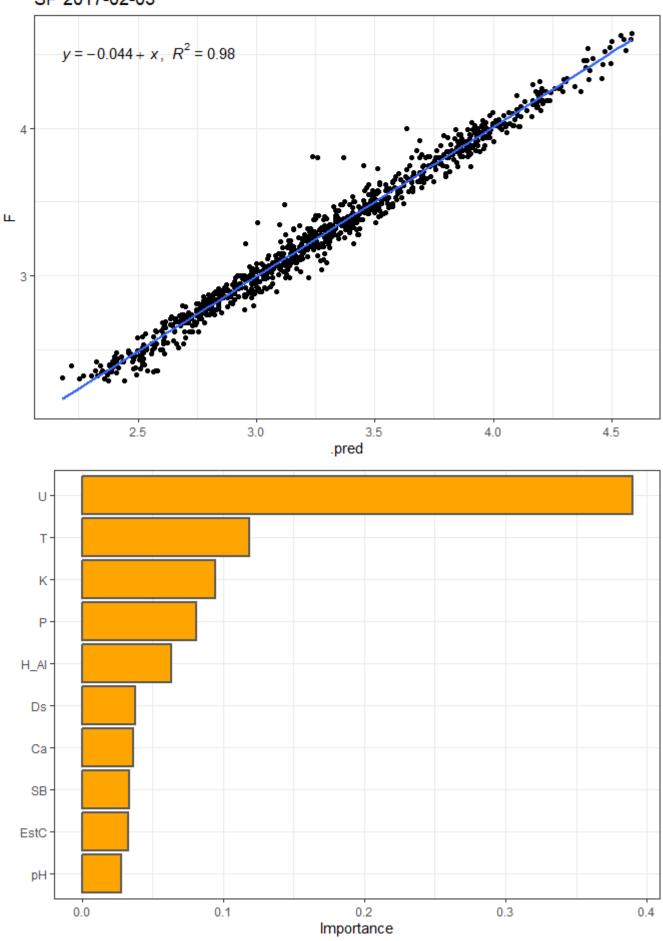


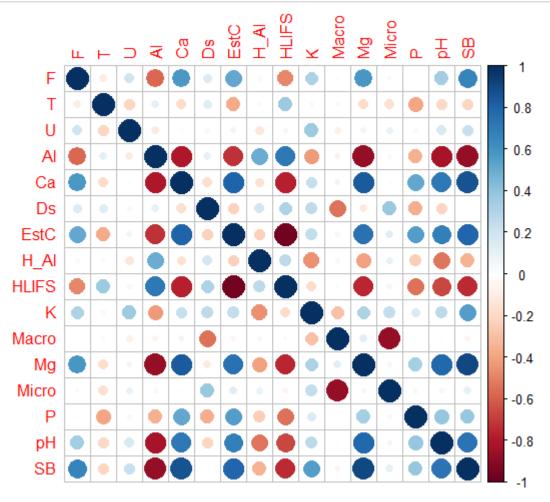


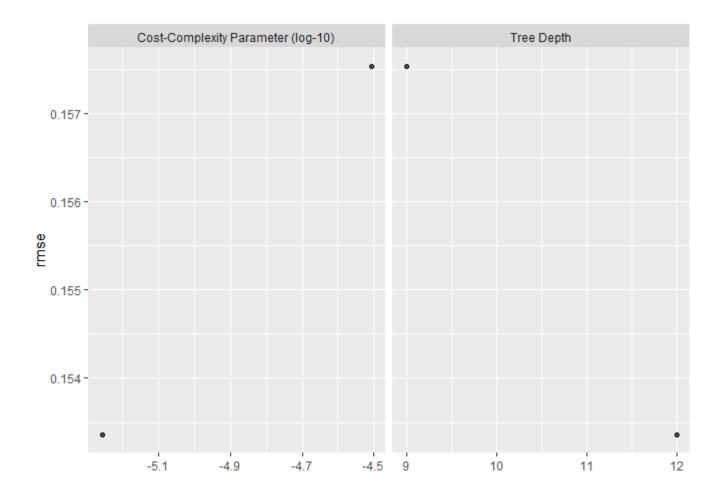


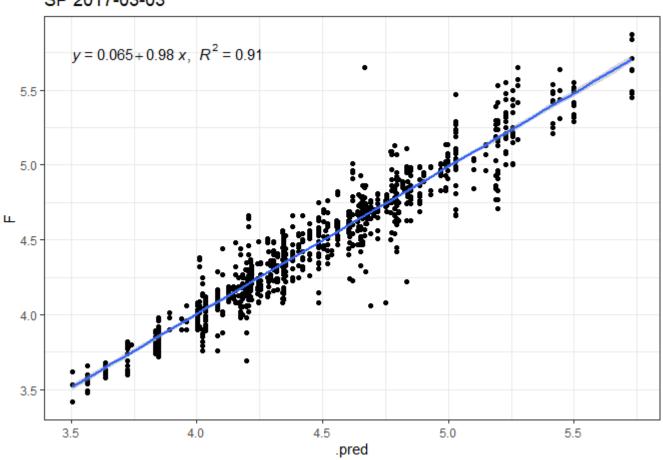


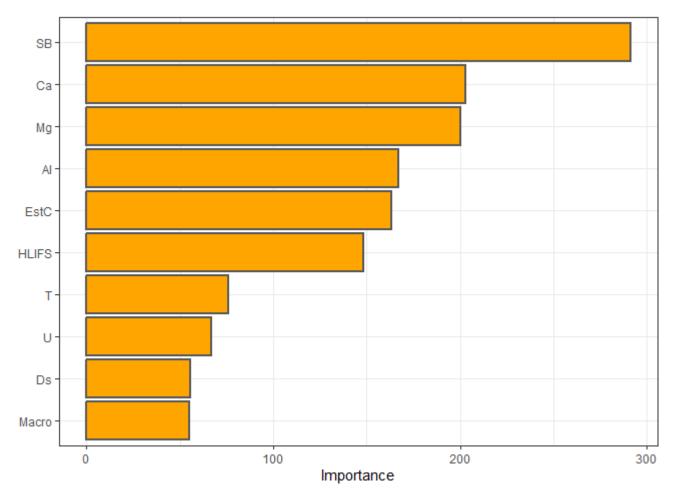


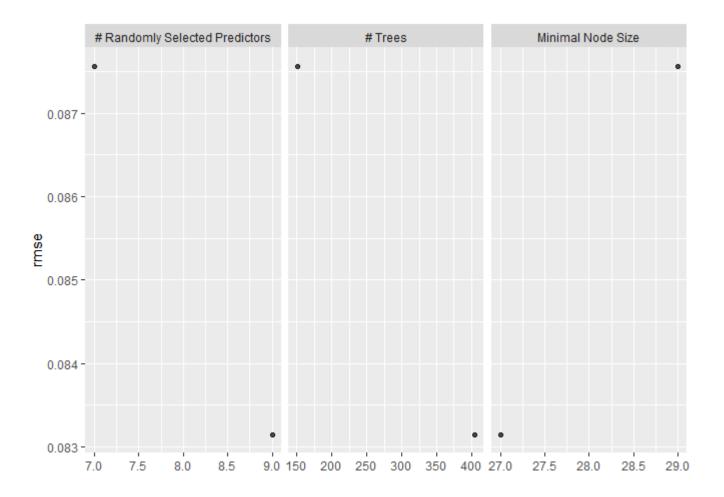


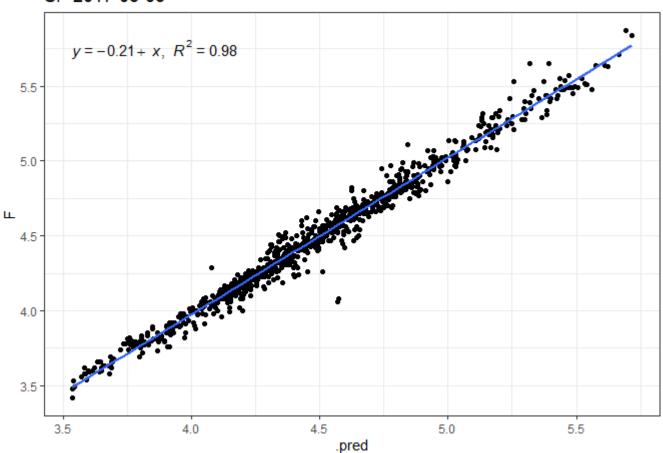


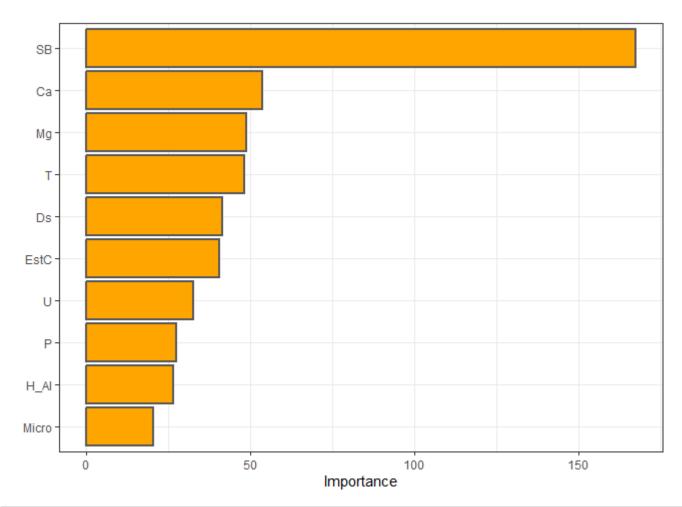


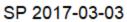


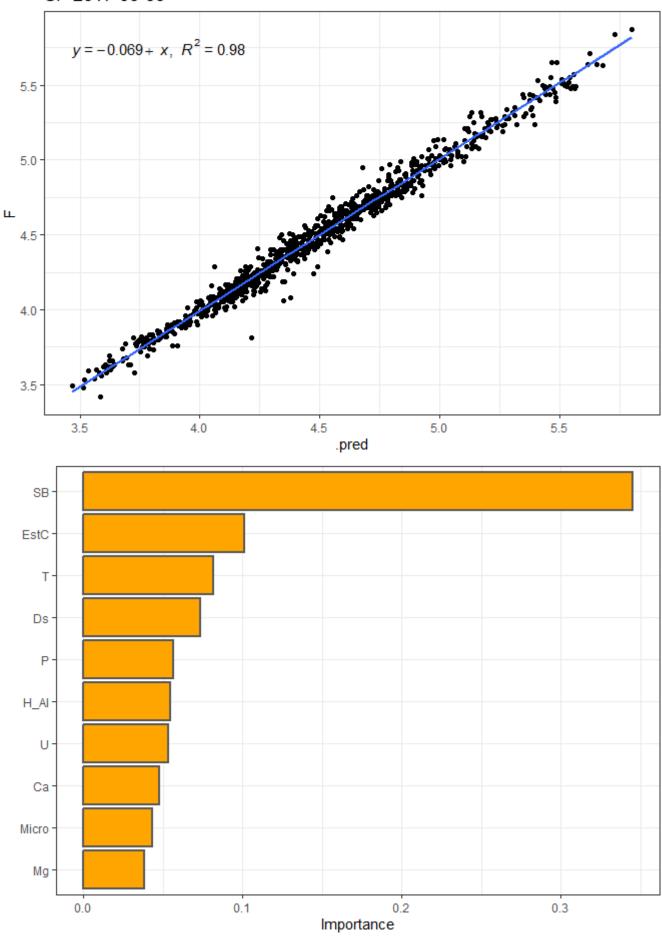


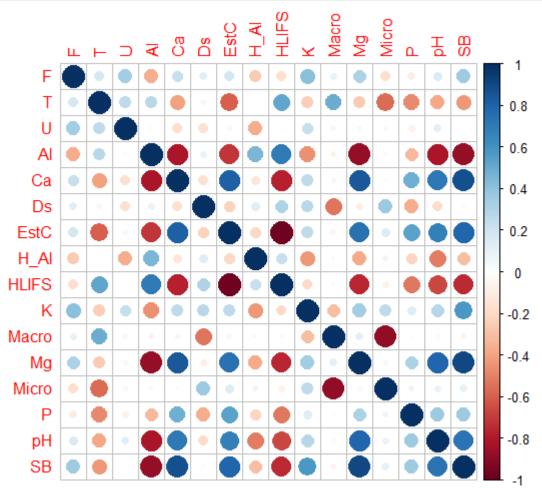


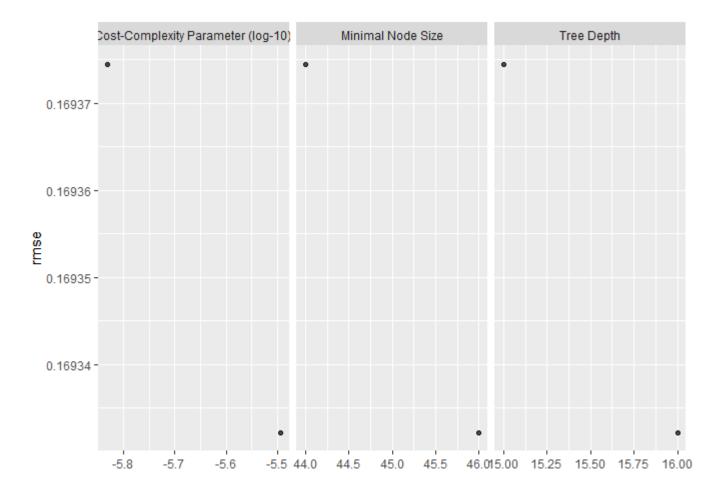


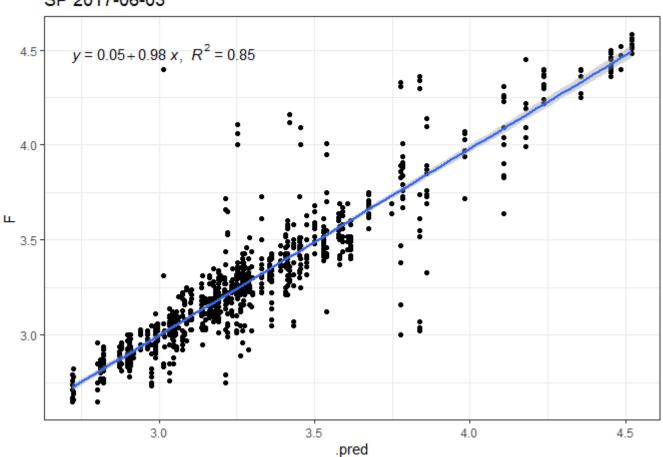


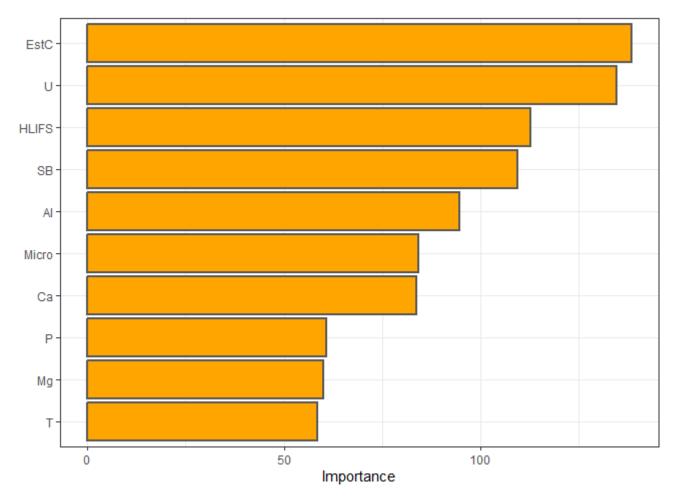


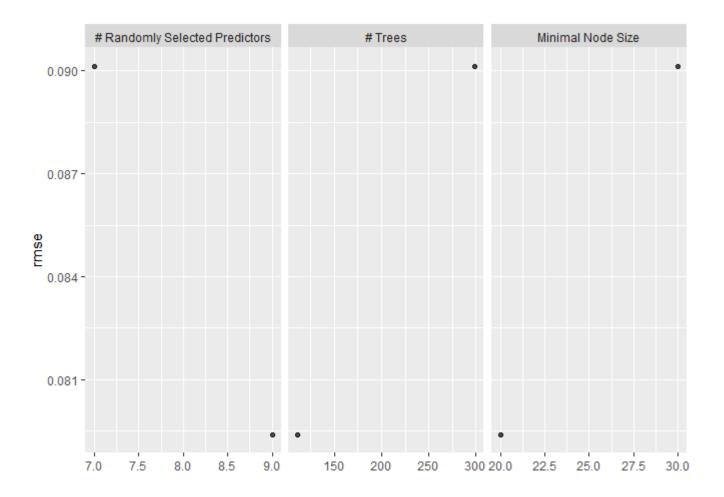


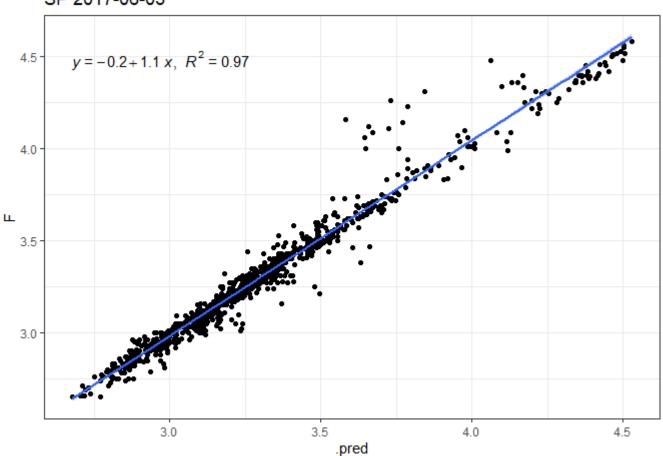


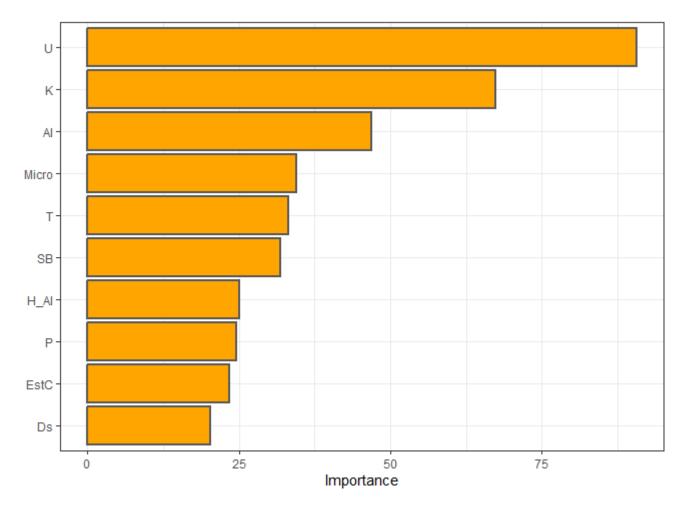


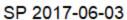


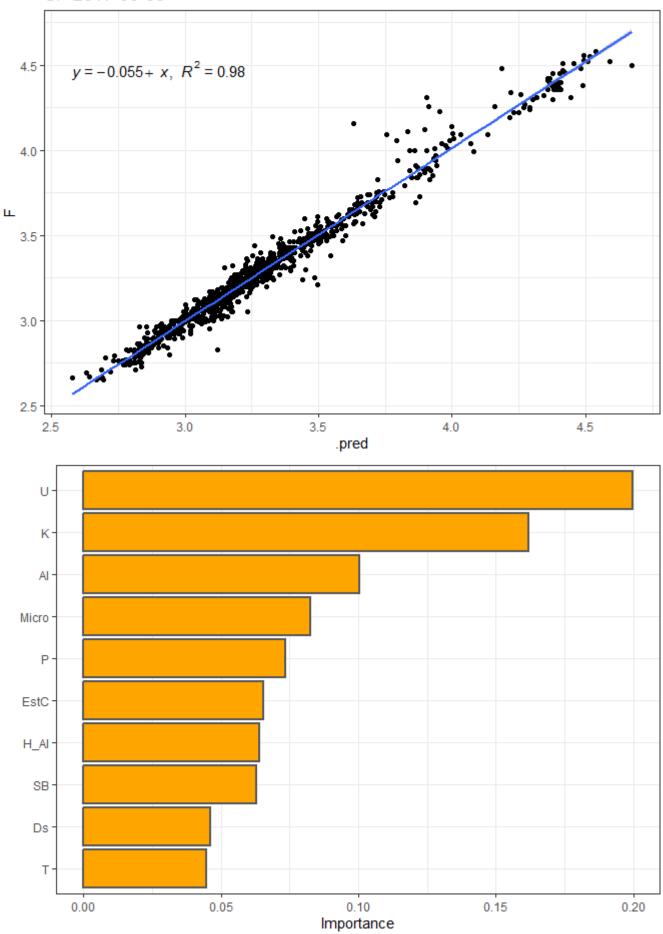


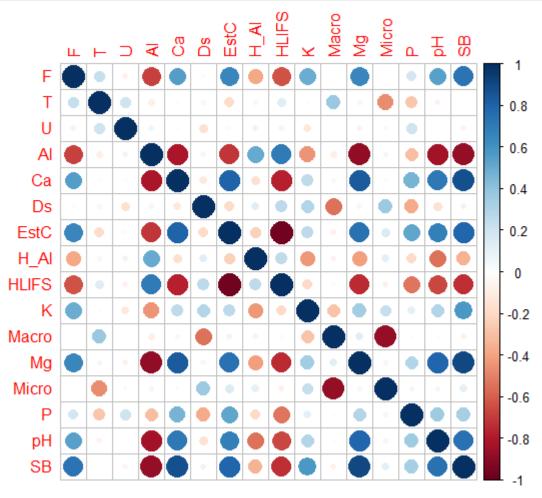


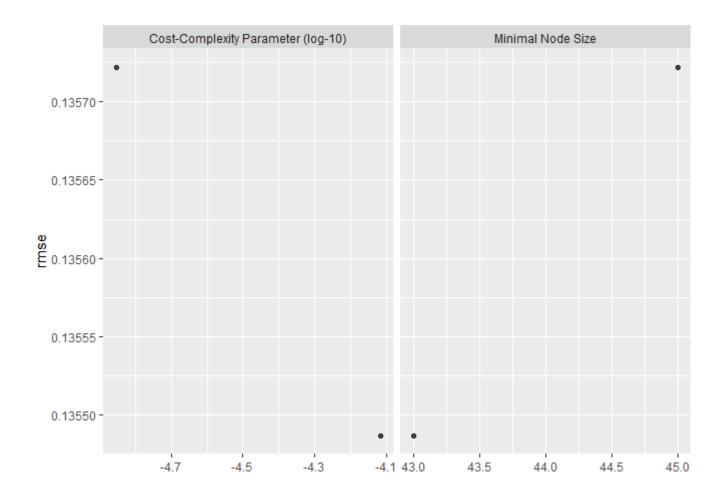


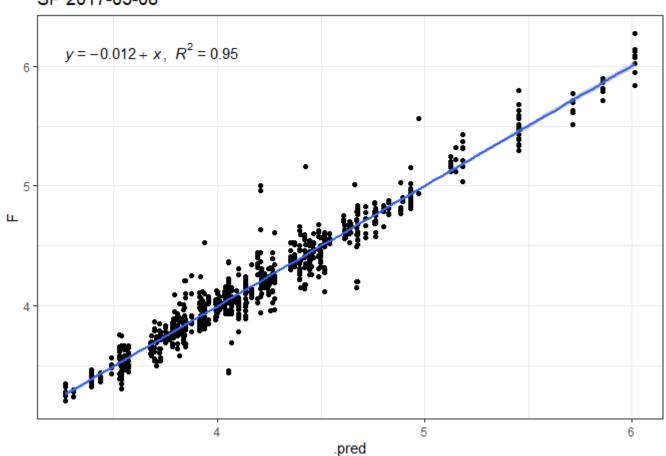


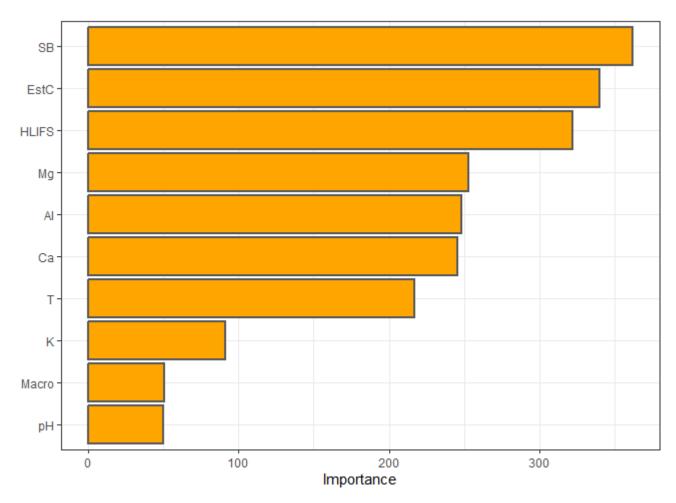


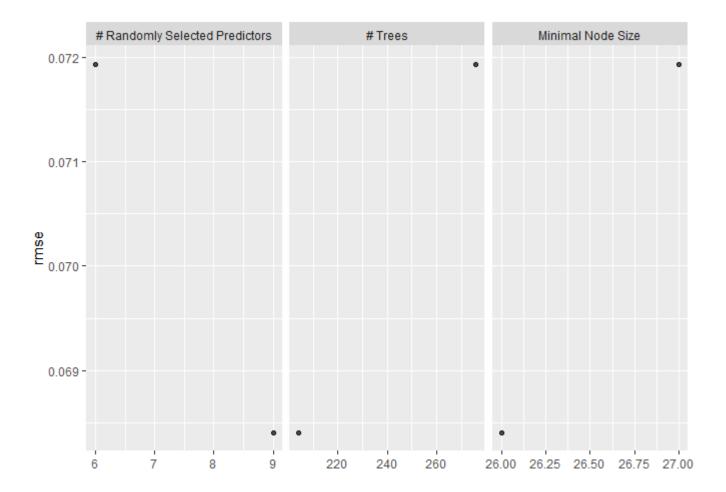


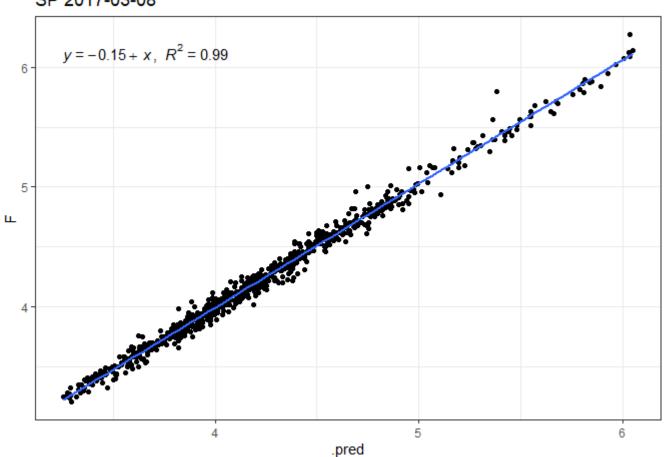


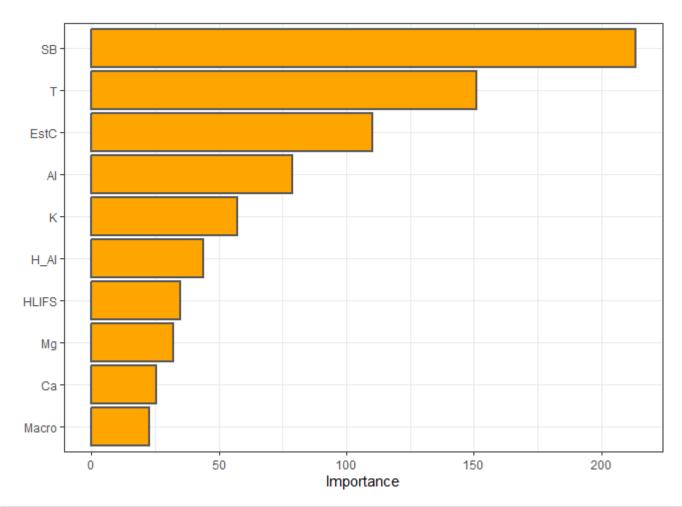




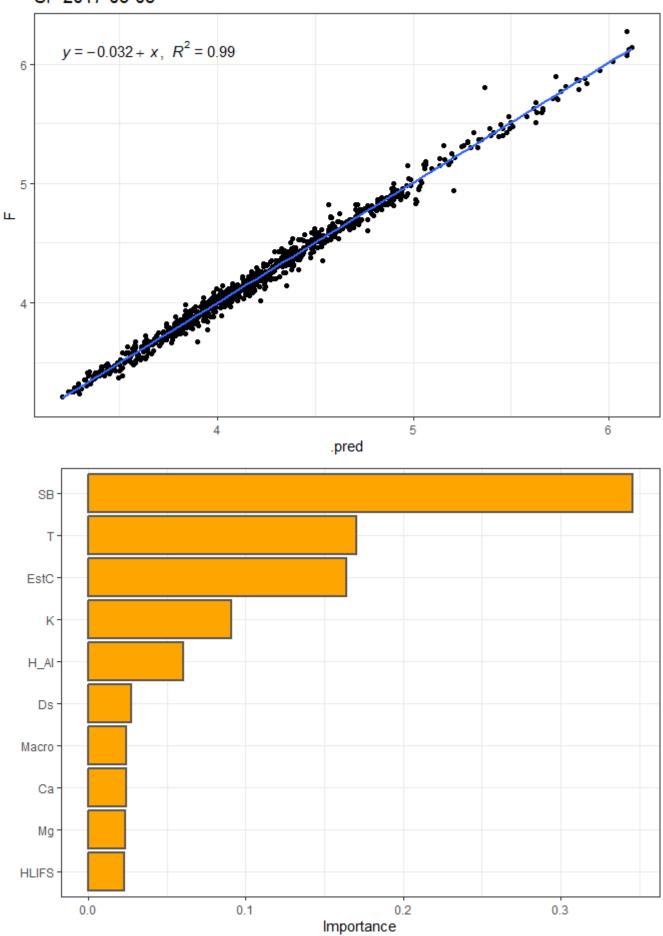


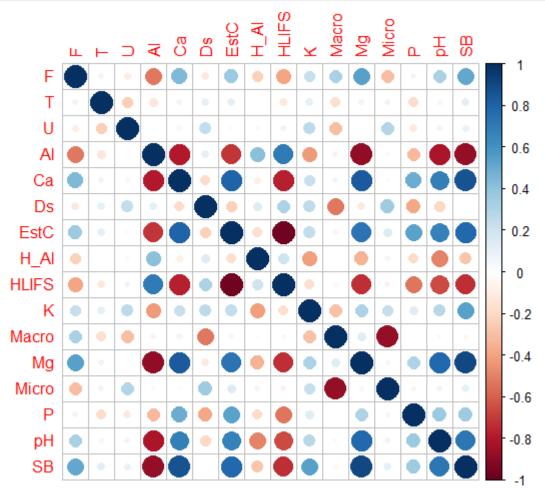


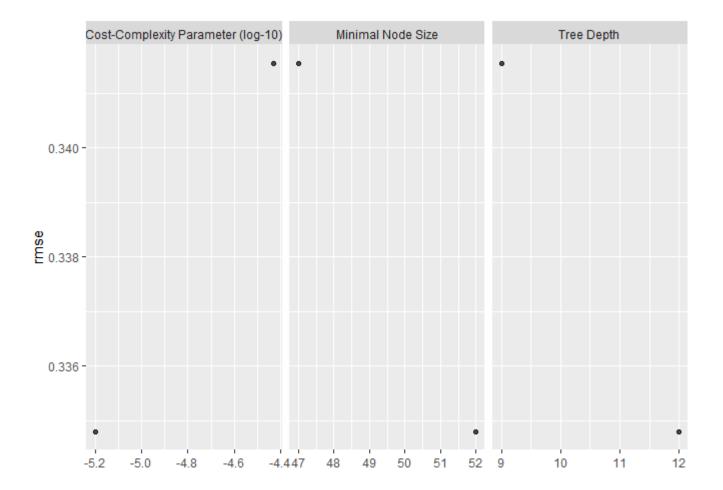


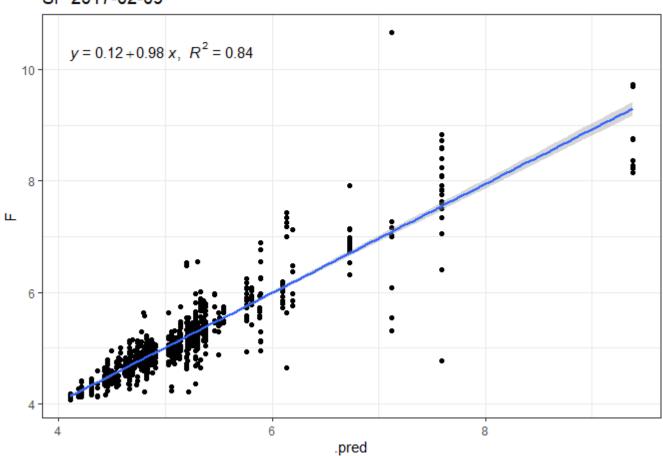


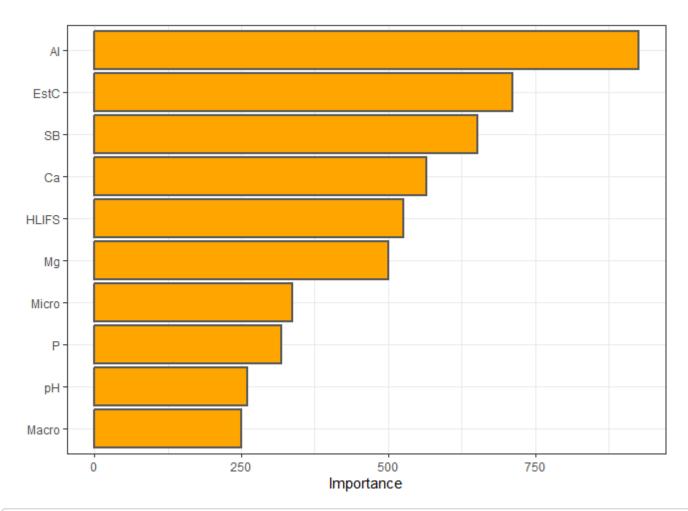


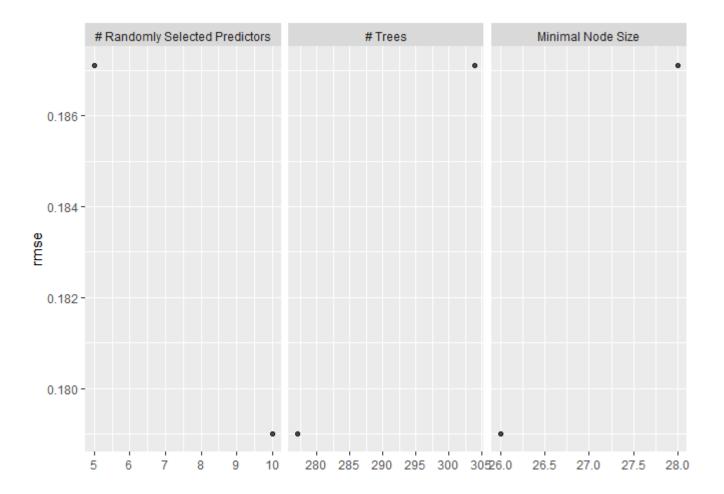


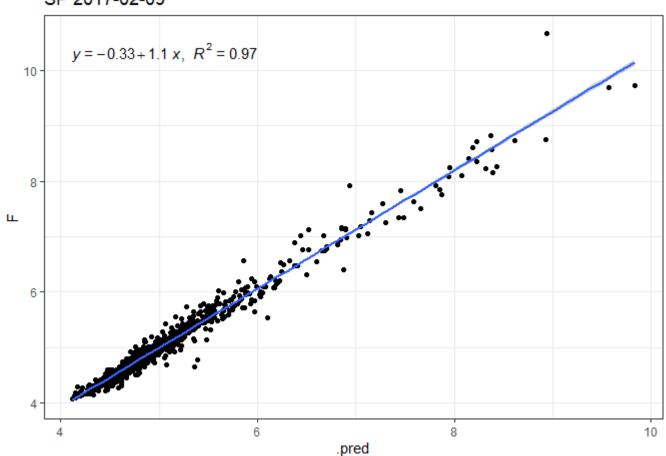


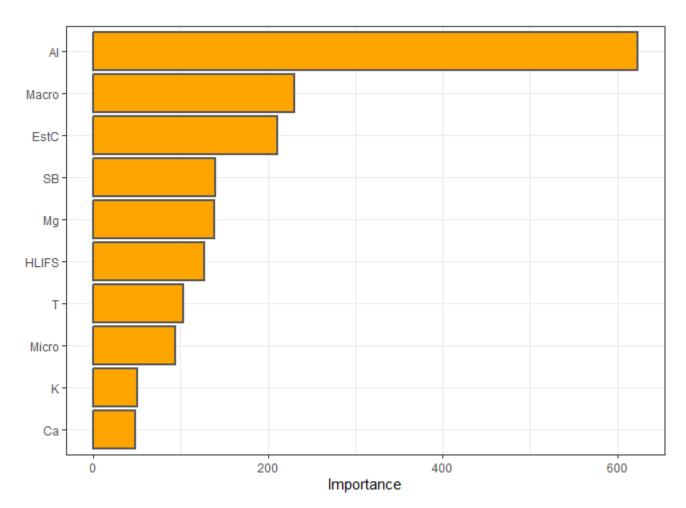


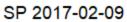


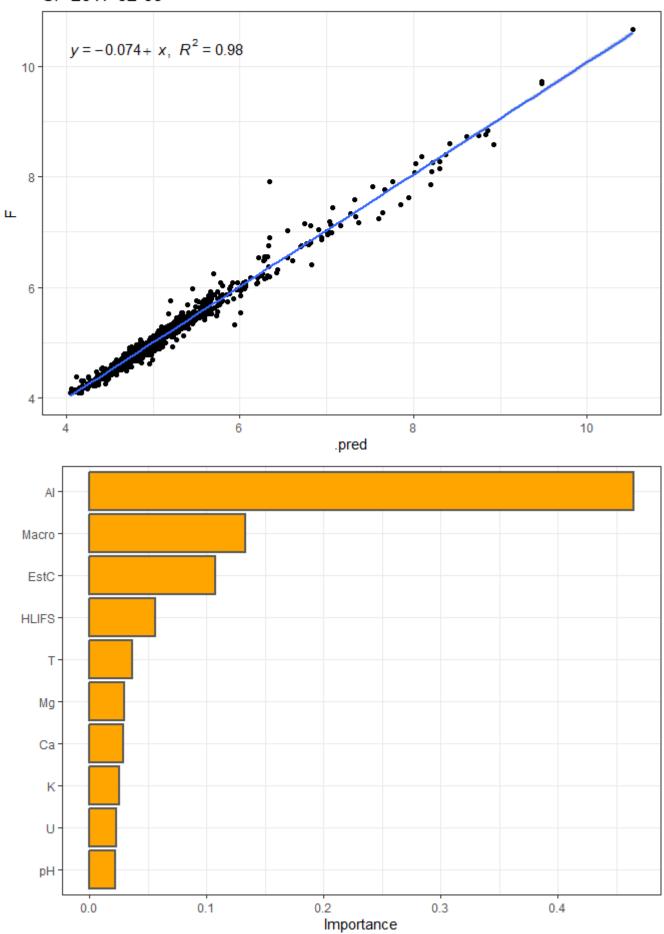


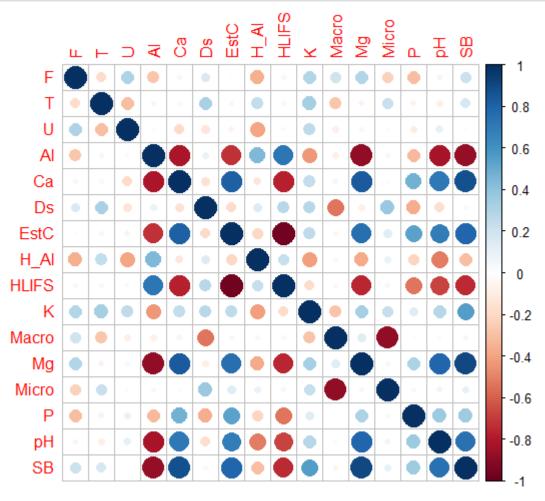


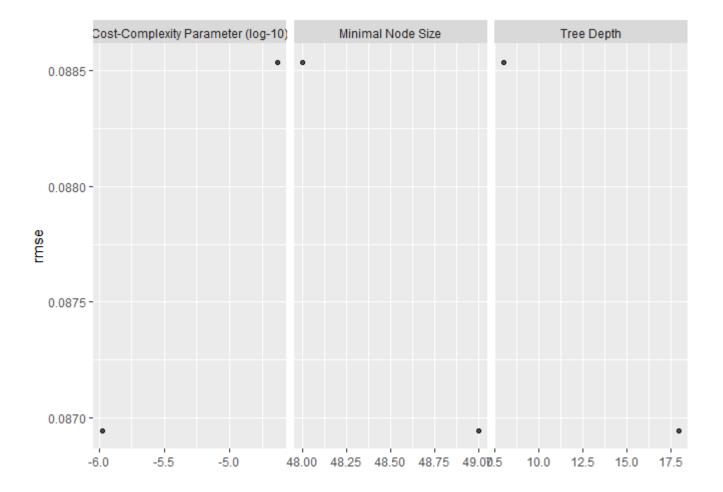




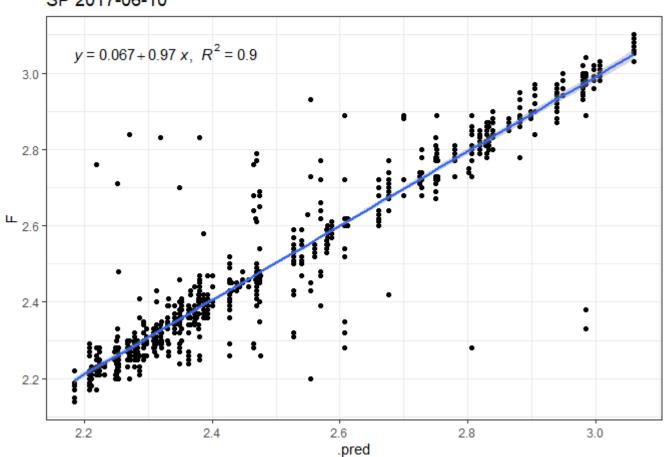


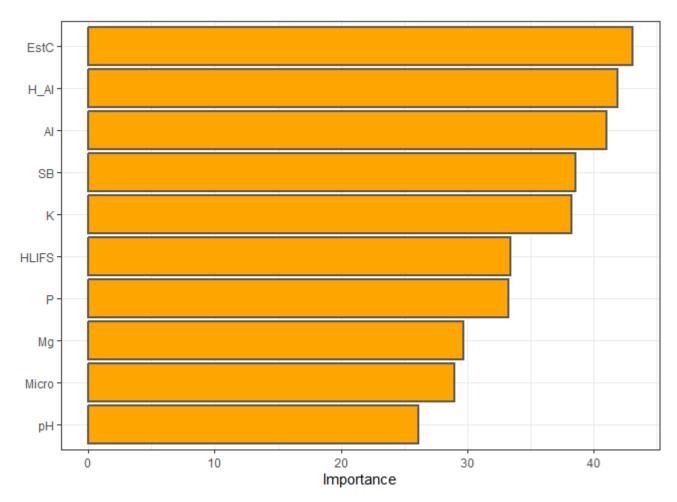


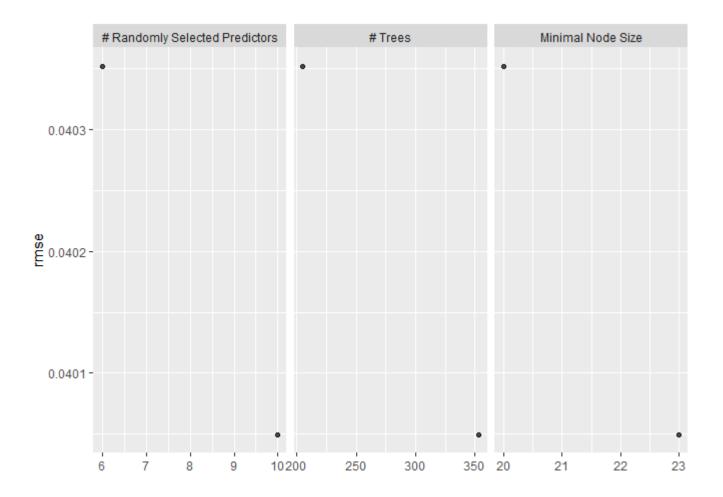




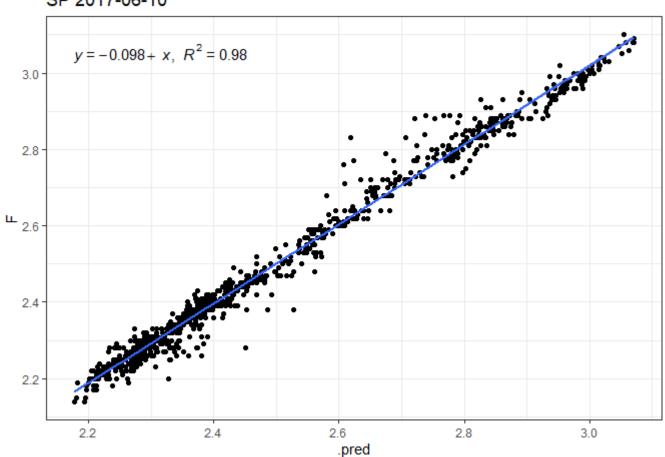
SP 2017-06-10

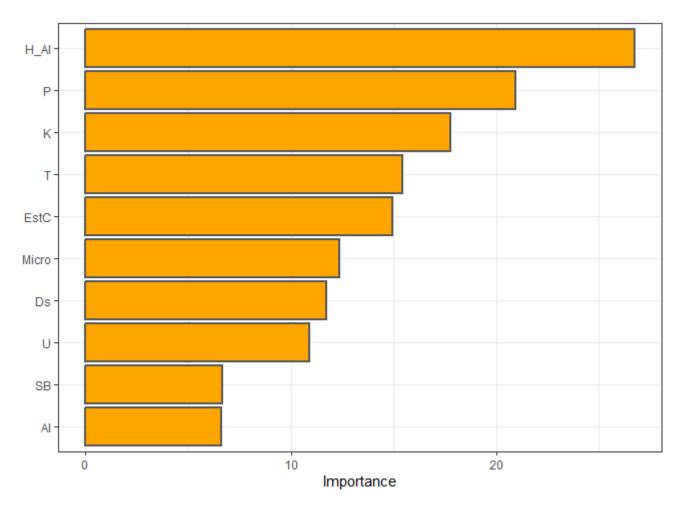


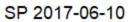


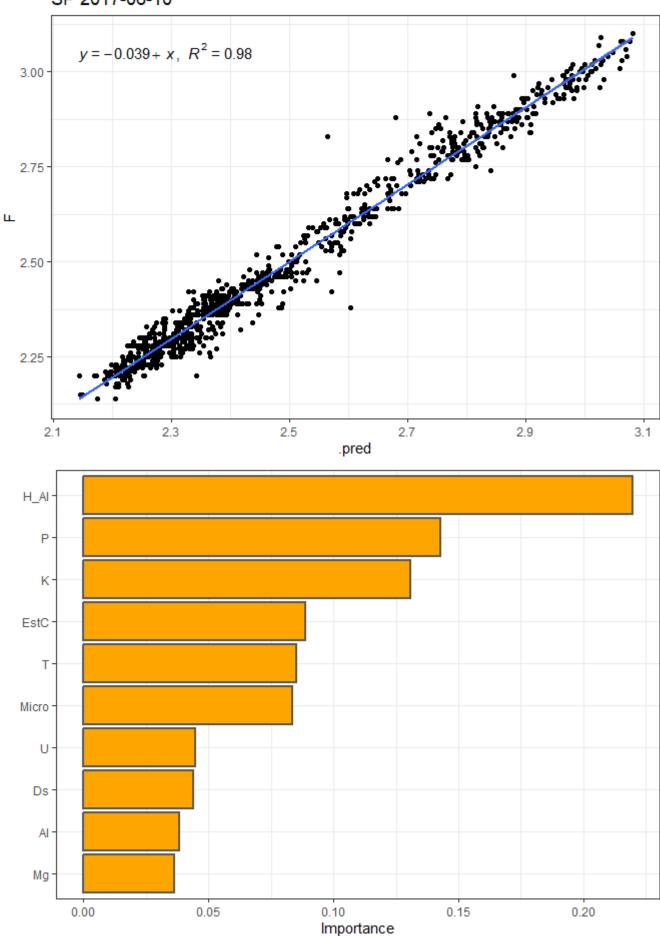


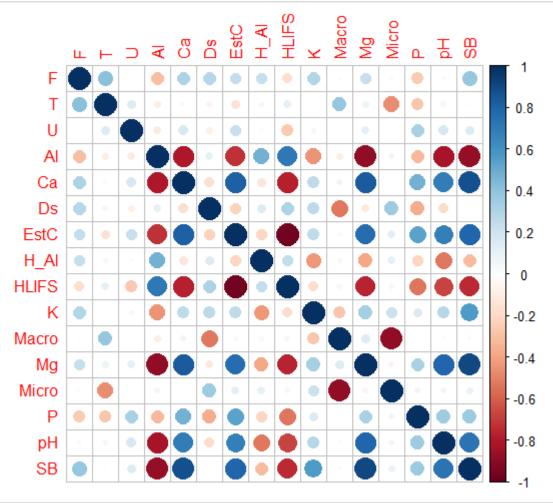
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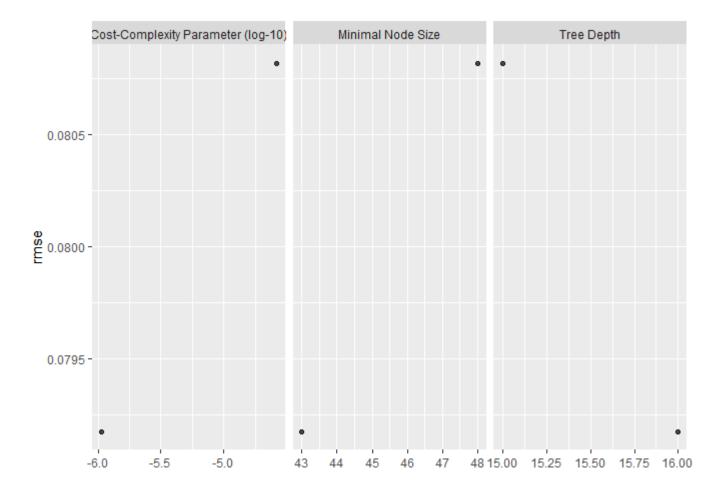


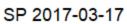


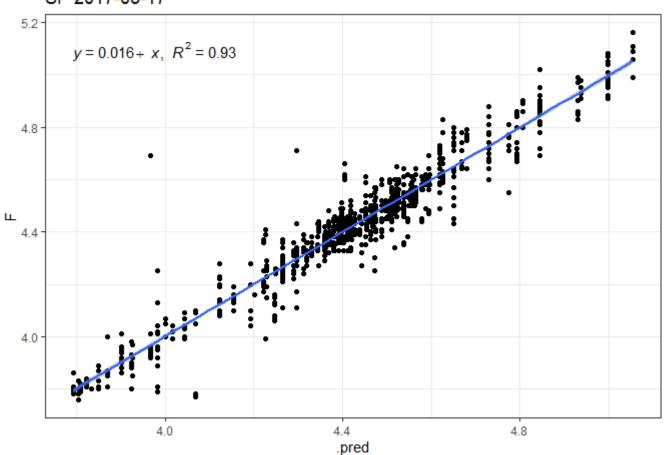


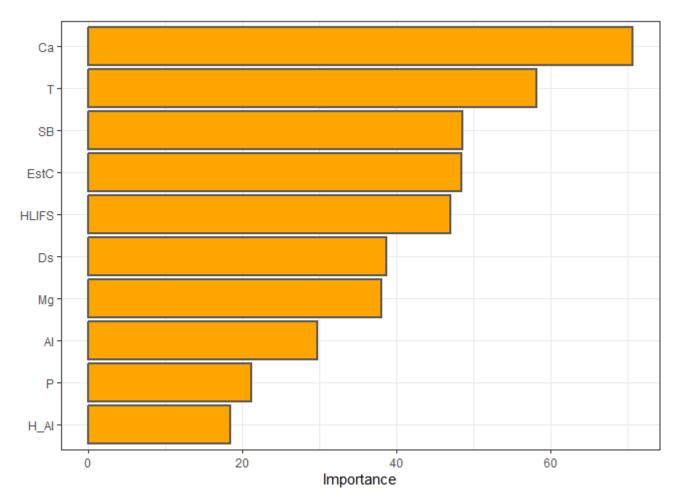


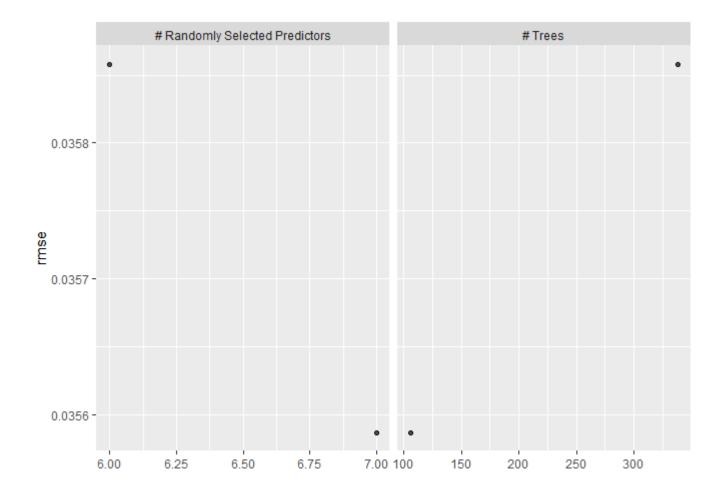


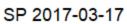


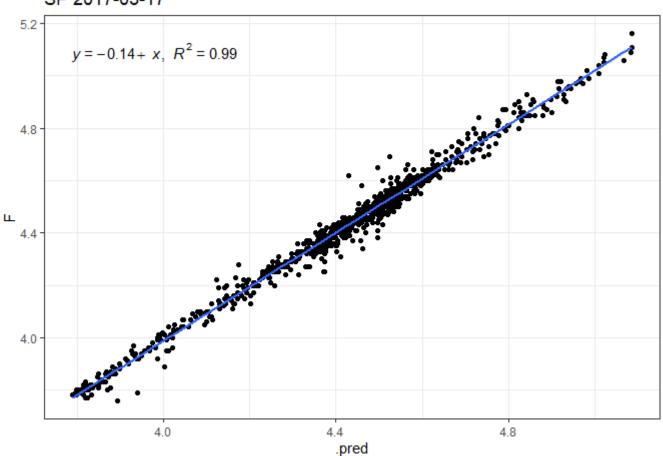


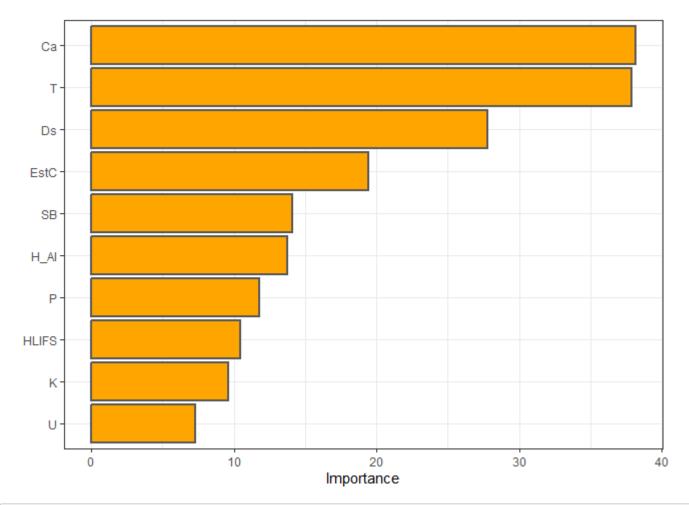


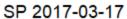


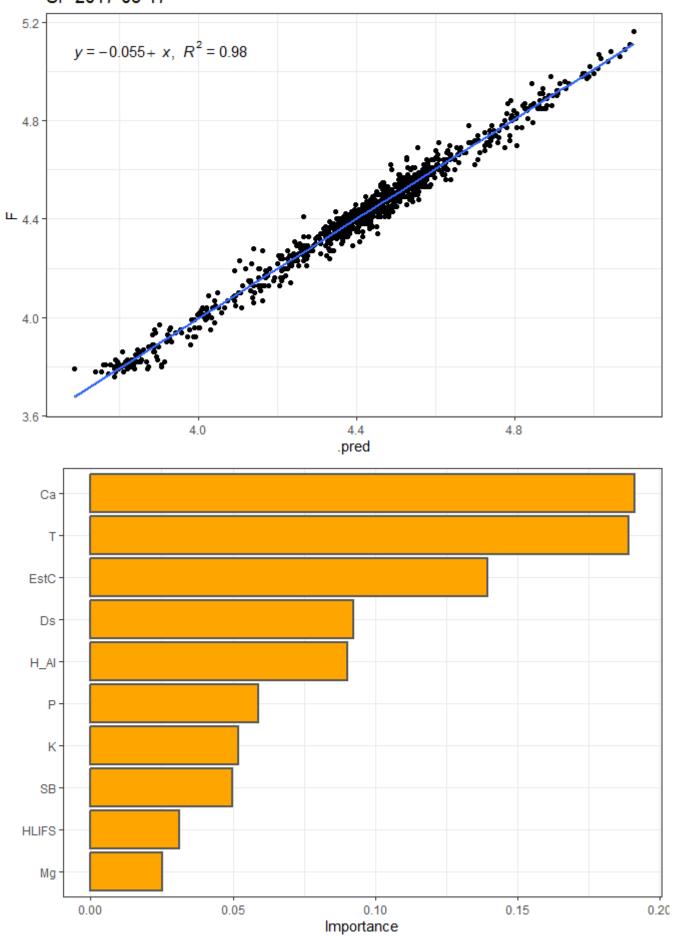


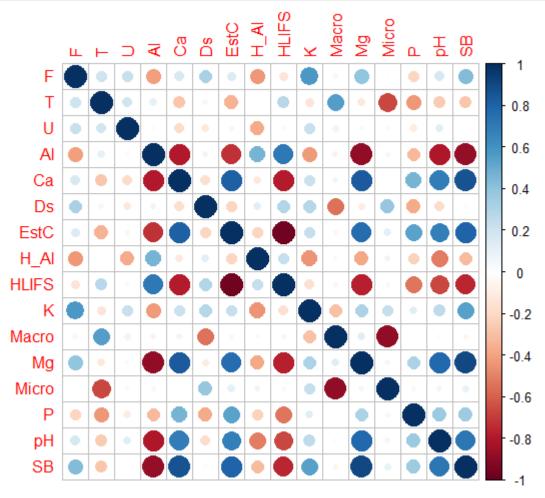


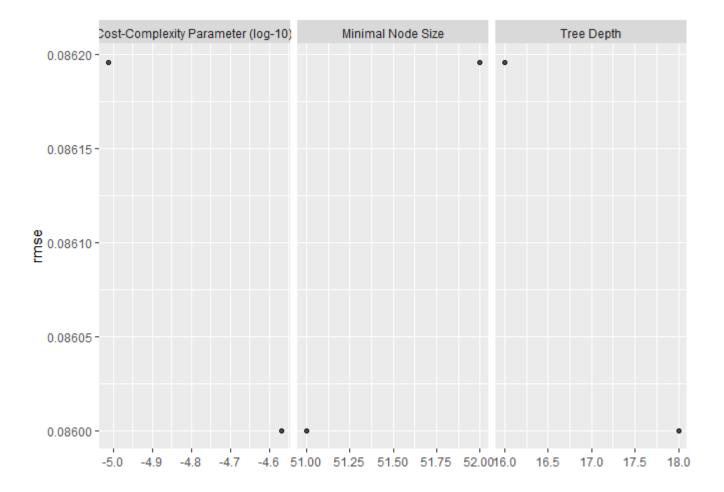


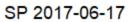


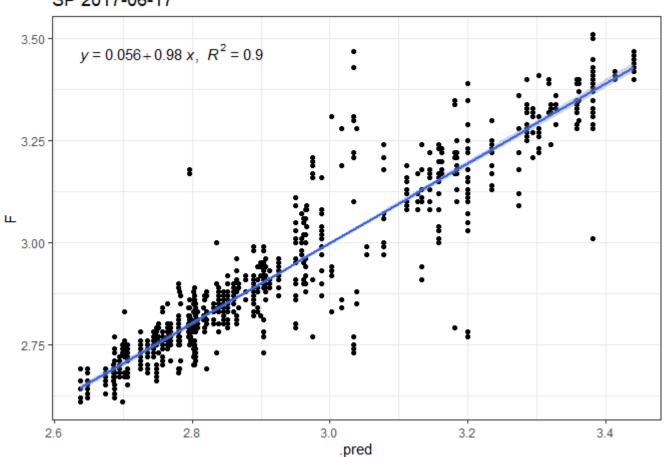


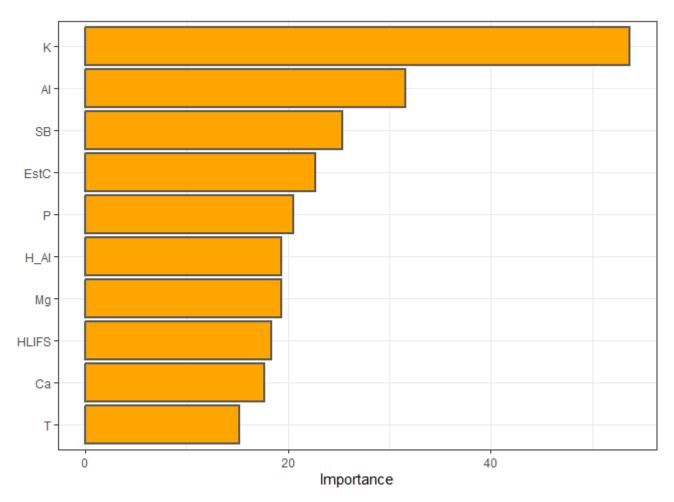


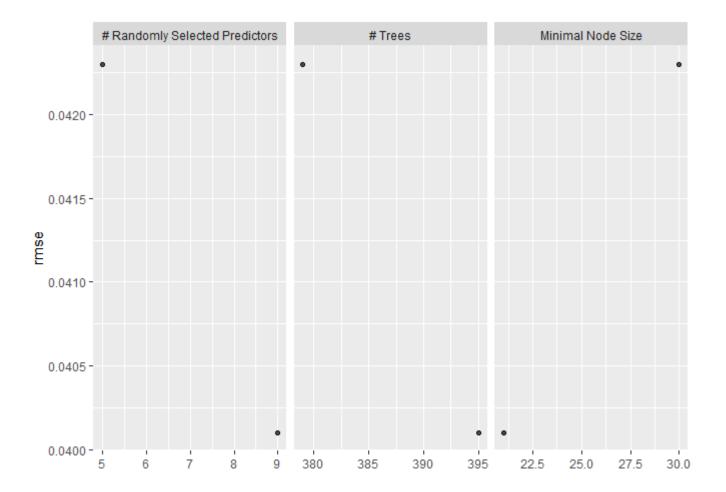




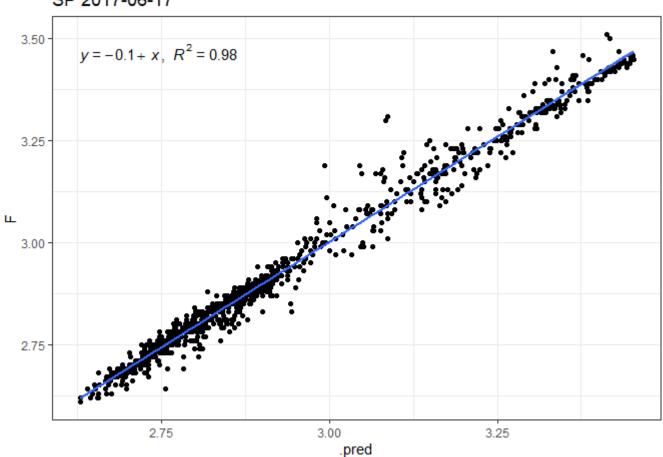


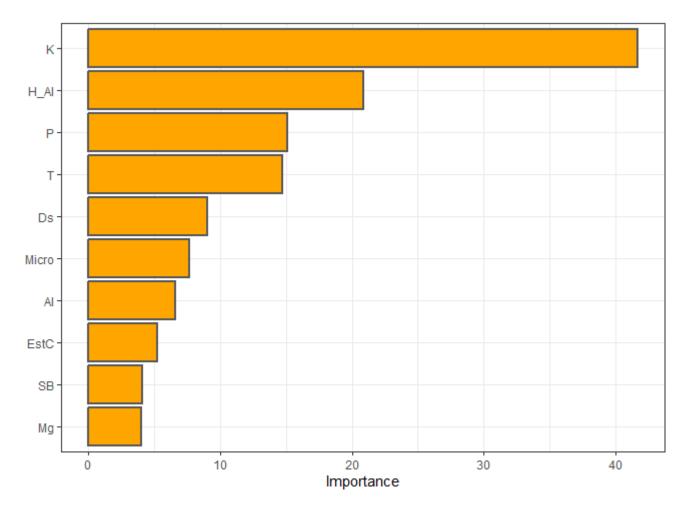




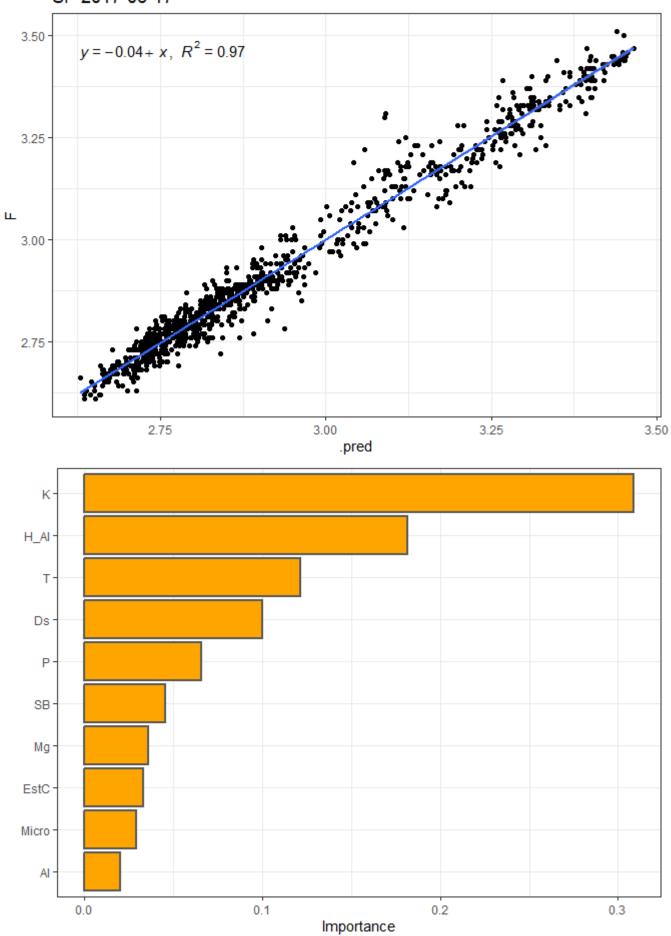


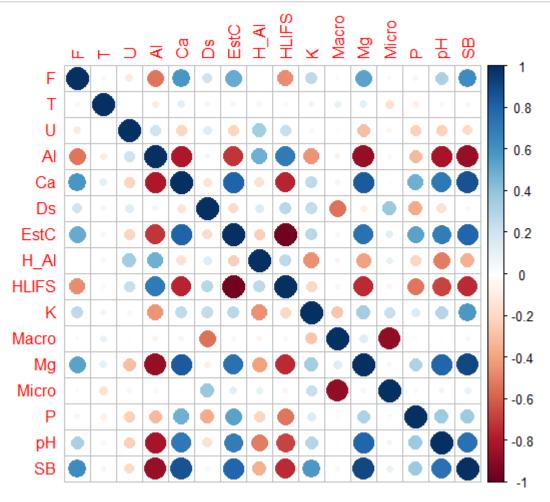


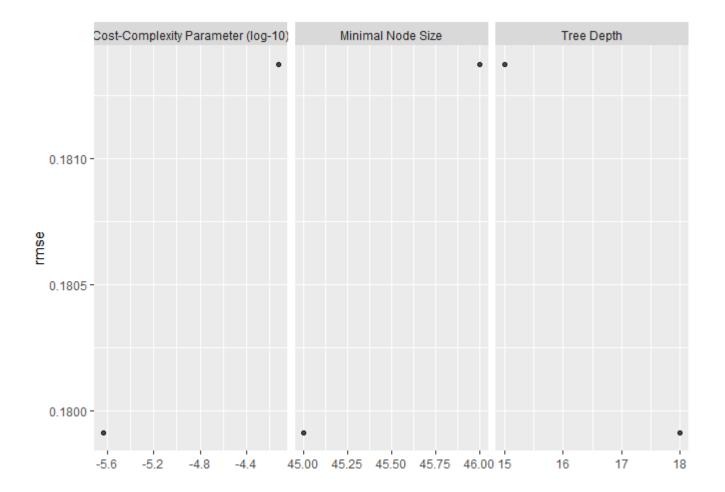




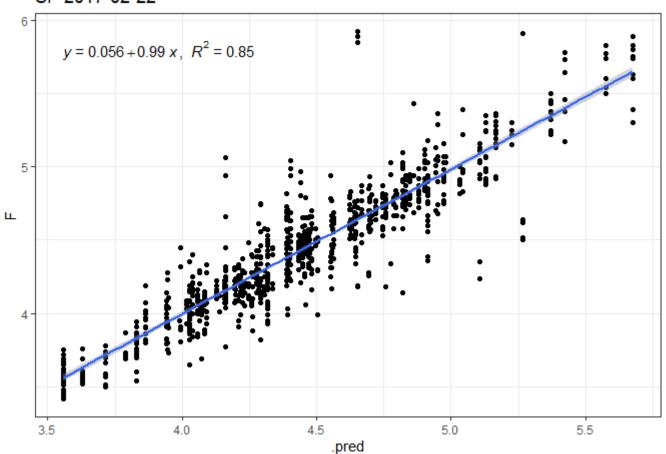


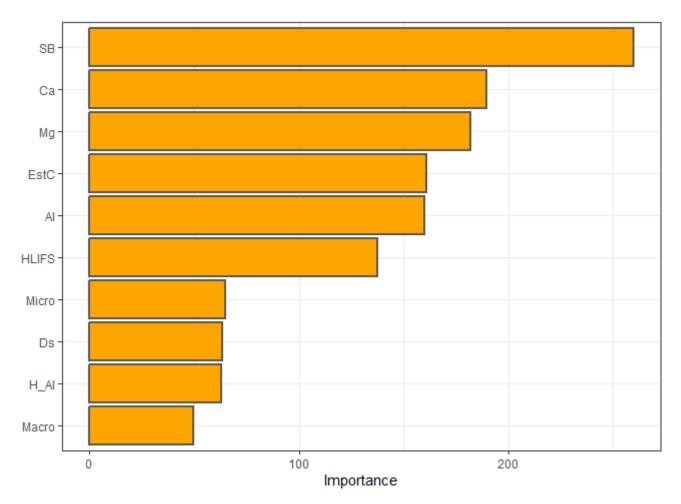


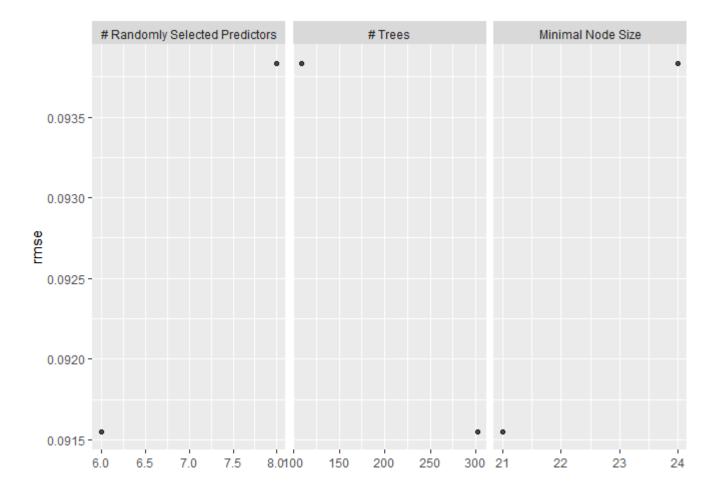


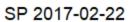


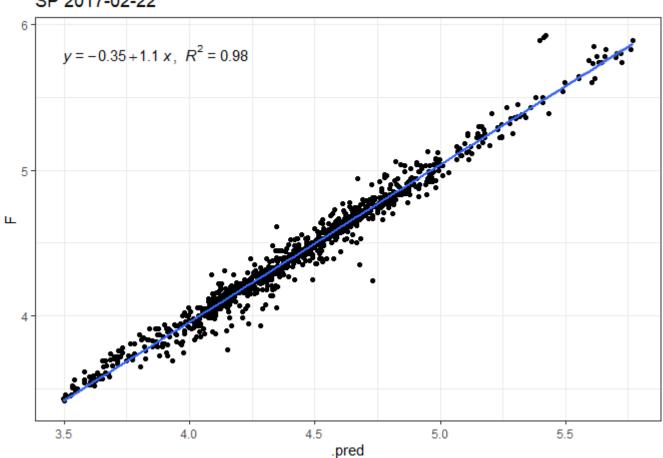
SP 2017-02-22

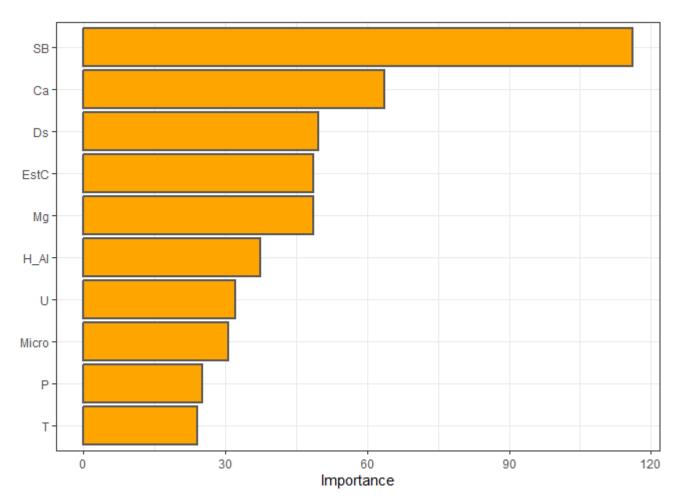


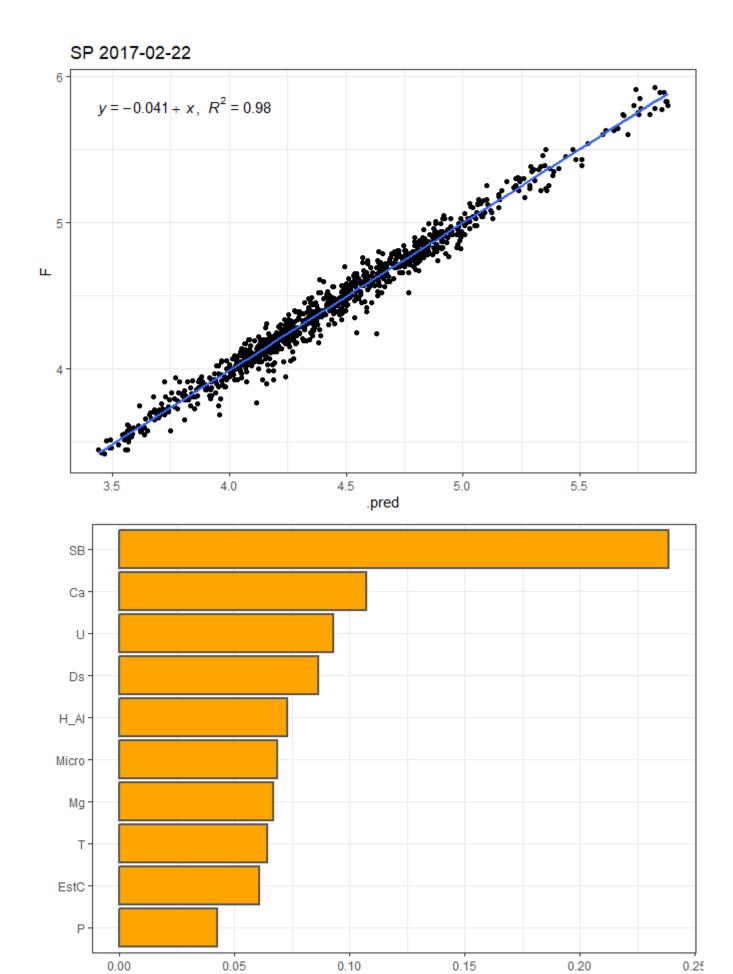












Importance

Material de Apêndice - Sinais (mapas dos atributos geoespacializados)

Mapas Eucalipto

```
# for(i in seq(files_eu)) {
# mp<-read.table(files_eu[i],skip = 5)
# image(mp %>% as.matrix(),xlab = files_eu[i])
# }
```

Mapas Silvipastoril

```
# for(i in seq(files_sp)) {
# mp<-read.table(files_sp[i],skip = 5)
# image(mp %>% as.matrix(),xlab = files_sp[i])
# }
```