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)

```
temporal_eu <- eu %>%
  filter(str_detect(nome, "^[F|U|T]")) %>%
  mutate(numero = as.numeric(str_remove(nome, "F|T|U")),
        ano = numero %% 10000,
        mes = numero %% 1000000 %/% 10000,
        dia = numero %/% 1e6,
        nome = str_remove_all(nome, "[0-9]")) %>%
  pivot_wider(names_from = nome,
        values_from = vetor)
```

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)
```

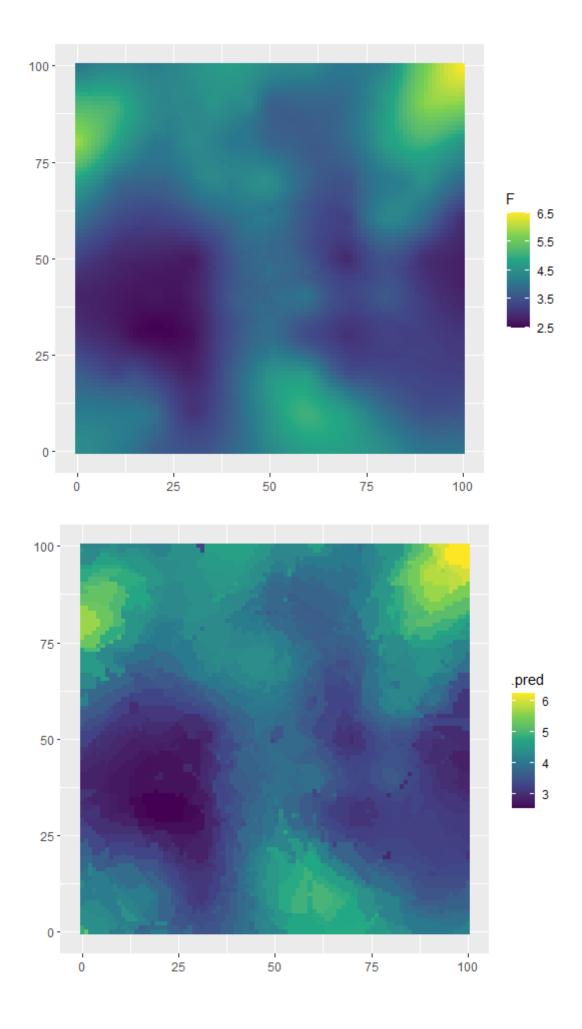
```
tibble(xnames=names(data_eu), ynames=names(data_sp)) %>%
 mutate(logic test = xnames == ynames)
#> # A tibble: 21 x 3
     xnames ynames logic_test
#> <chr> <chr> <lgl>
#> 1 id id
                   TRUE
#> 2 data data TRUE
#> 3 ano ano TRUE
\#> 4 mes mes TRUE
#> 5 dia dia TRUE
#> 6 F F TRUE
#> 7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

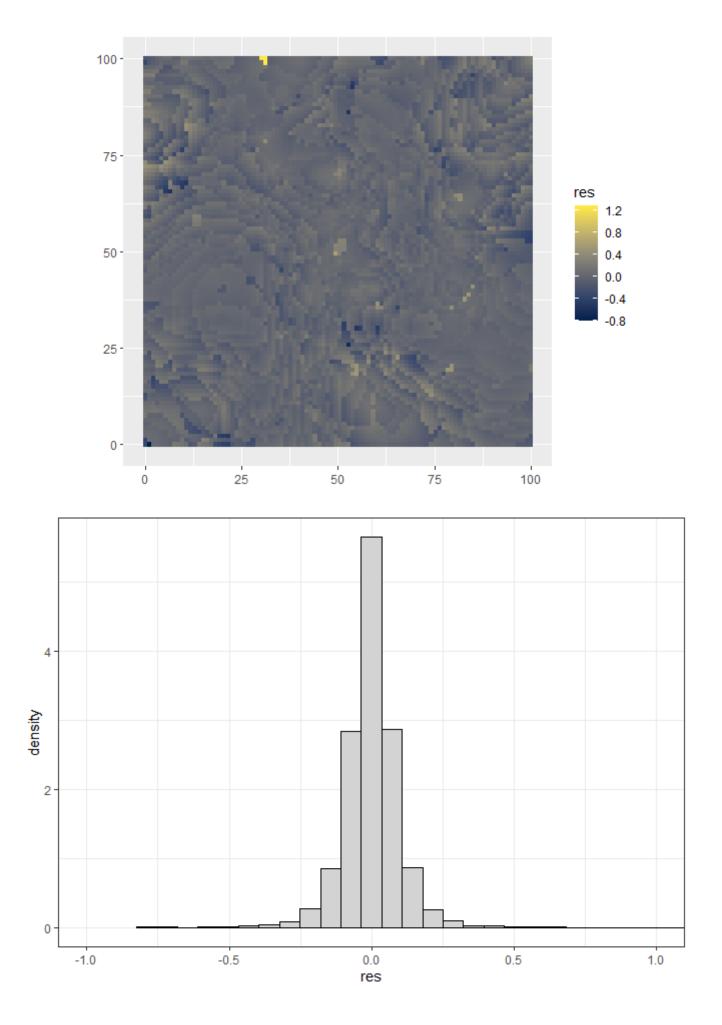
Unindo toda a base de dados

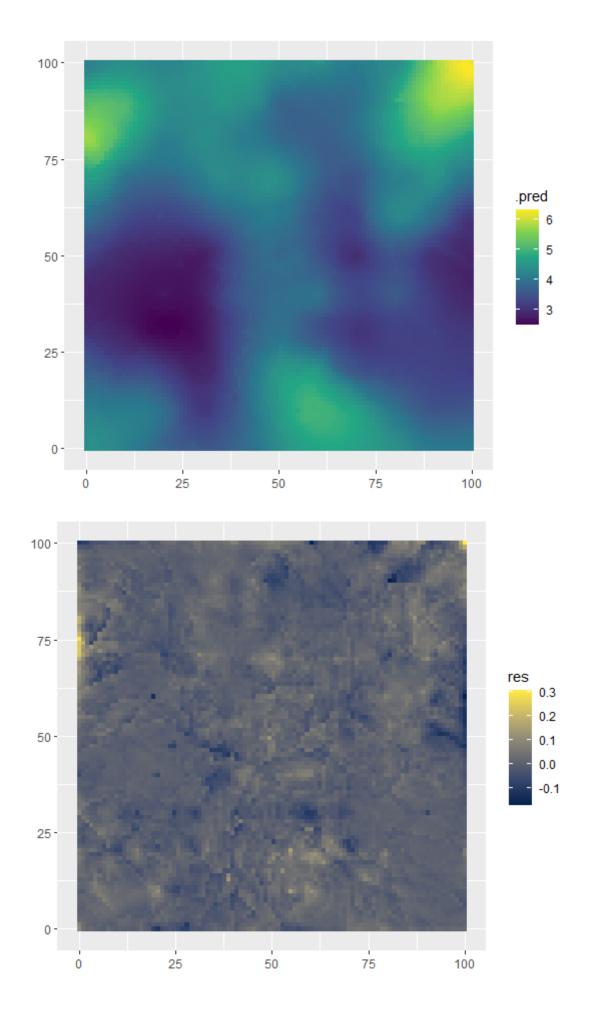
Criando os preditos para cada dia - EUCALIPTO

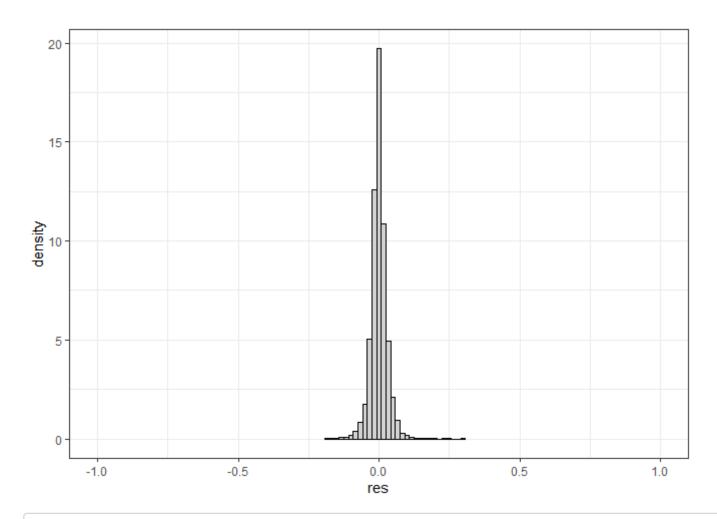
```
dias <- data eu$data %>% unique()
dis <- 100/93
grid <- expand.grid(Y=seq(0,100,dis), X=seq(0,100,dis))</pre>
for(i in seq along(dias)){
 di <- dias[i]</pre>
 data eu %>% filter(data == di)
 file models eu <- list.files("models-3", pattern = "EU")</pre>
 file models eu <- grep(paste0(di), file models eu, value = TRUE)
 fco2 modelo load dt <- read rds(
   paste0("models-3/",grep("dt",file models eu,value=TRUE)
   ))
 fco2_modelo_load_rf <- read_rds(</pre>
   paste0("models-3/",grep("rf",file models eu,value=TRUE)
 fco2 modelo load xgb <- read rds(
   paste0("models-3/",grep("xgb",file models eu,value=TRUE)
 yp dt <- predict(fco2 modelo load dt, new data = data eu %>%
                     filter(data == di))
 yp_rf <- predict(fco2_modelo_load_rf, new_data = data_eu %>%
                     filter(data == di))
  # yp xgb <- predict(fco2 modelo load xgb, new data = data eu %>%
                  filter(data == di))
 mp krg <- tibble(grid, data eu %>%
                     filter(data == di)) %>%
   ggplot(aes(x=X,y=Y)) +
   geom tile(aes(fill = F)) +
   scale fill viridis c() +
   coord_equal()+labs(x="",y="")
 mp dt <- tibble(grid, data eu %>%
                    filter(data == di), yp dt) %>%
   ggplot(aes(x=X,y=Y)) +
   geom tile(aes(fill = .pred)) +
   scale fill viridis c() +
   coord equal() +labs(x="",y="")
 mp rf <- tibble(grid, data eu %>%
                    filter(data == di), yp rf) %>%
   ggplot(aes(x=X,y=Y)) +
   geom tile(aes(fill = .pred)) +
   scale fill viridis c() +
   coord equal() +labs(x="",y="")
 mp dt res <- tibble(grid, data eu %>%
```

```
filter(data == di), yp dt) %>%
   mutate(res = F - .pred) %>%
   ggplot(aes(x=X,y=Y)) +
   geom tile(aes(fill = res)) +
   scale_fill_viridis_c(option = "E") +
   coord equal() +labs(x="",y="")
 hist_dt_res <- tibble(grid, data_eu %>%
                          filter(data == di), yp dt) %>%
   mutate(res = F - .pred) %>%
   ggplot(aes(x=res, y=..density..)) +
   geom histogram(color="black",fill="lightgray") +theme bw()+
   coord cartesian(xlim=c(-1,1))
 mp rf res <- tibble(grid, data eu %>%
                    filter(data == di), yp rf) %>%
   mutate(res = F - .pred) %>%
   qqplot(aes(x=X,y=Y)) +
   geom_tile(aes(fill = res)) +
   scale fill viridis c(option = "E") +
   coord_equal() +labs(x="",y="")
 hist_rf_res <- tibble(grid, data_eu %>%
                          filter(data == di), yp rf) %>%
   mutate(res = F - .pred) %>%
   ggplot(aes(x=res, y=..density..)) +
   geom histogram(color="black",fill="lightgray") +theme bw()+
   coord cartesian(xlim=c(-1,1))
 print(di)
 print(mp krg)
 print(mp dt)
 print(mp dt res)
 print(hist dt res)
 print(mp rf)
 print(mp rf res)
 print(hist rf res)
#> [1] "2017-06-03"
```

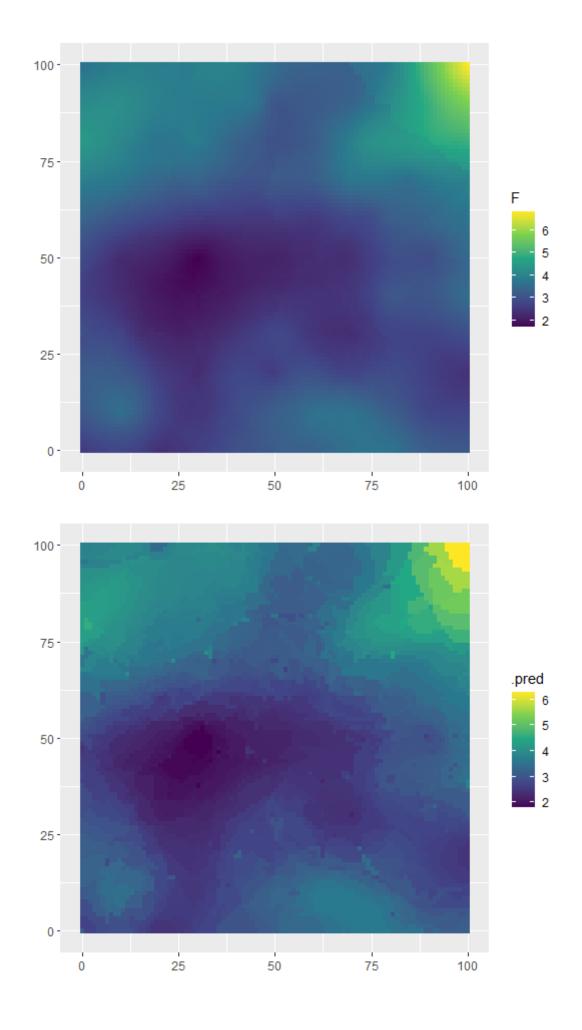


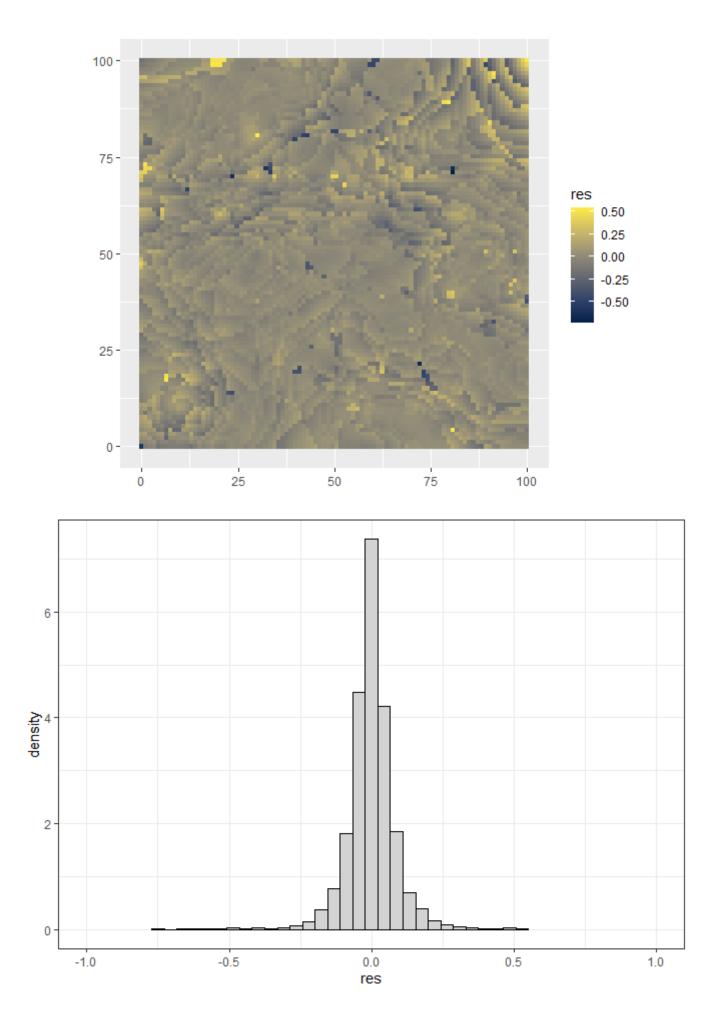


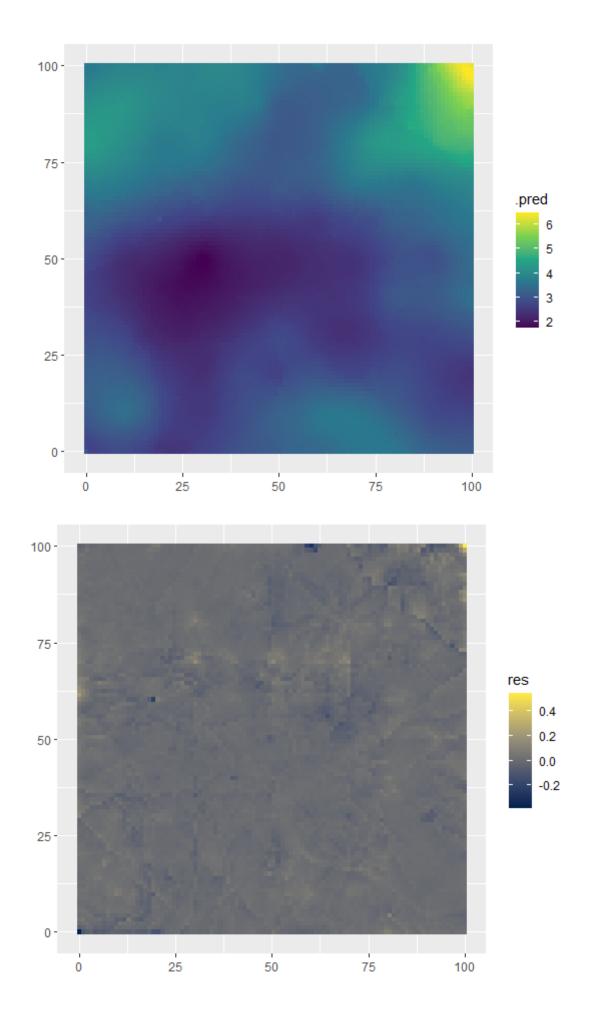


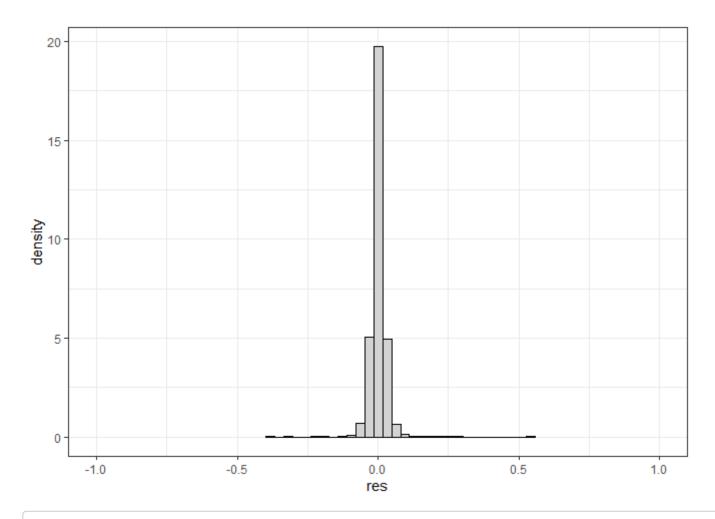


#> [1] "2017-06-10"

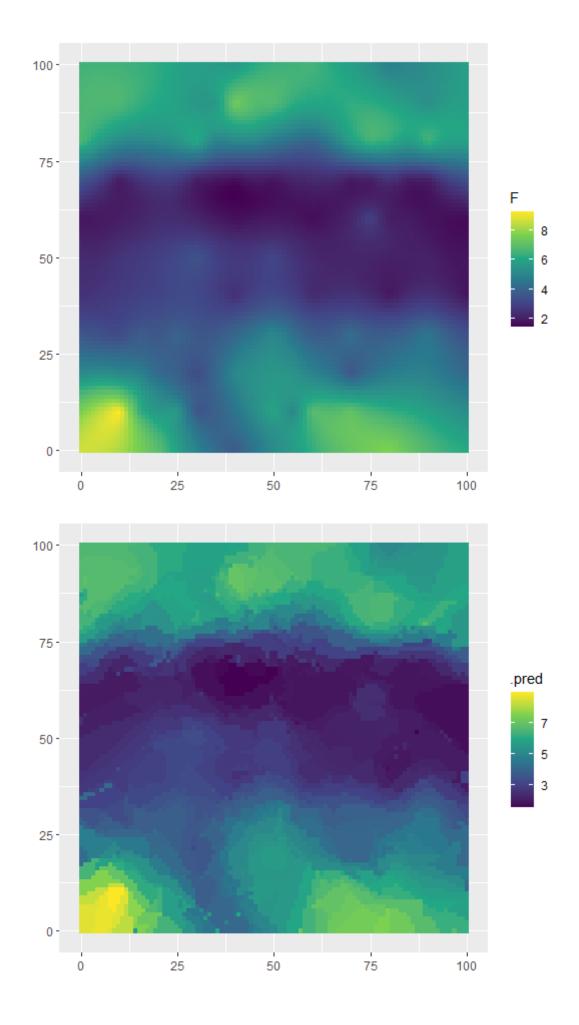


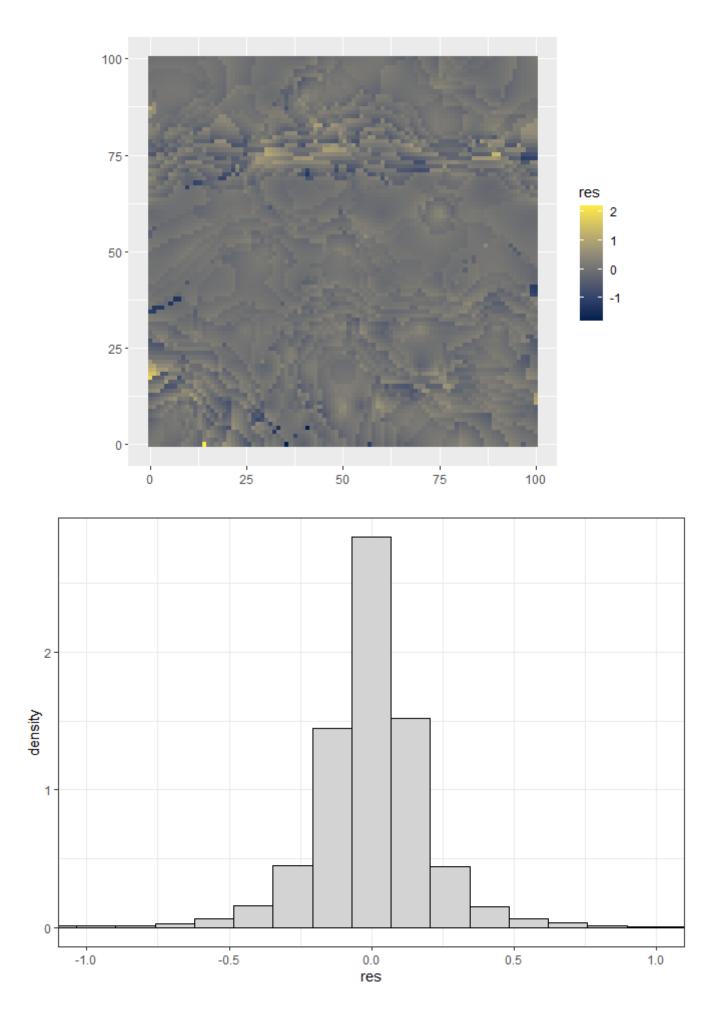


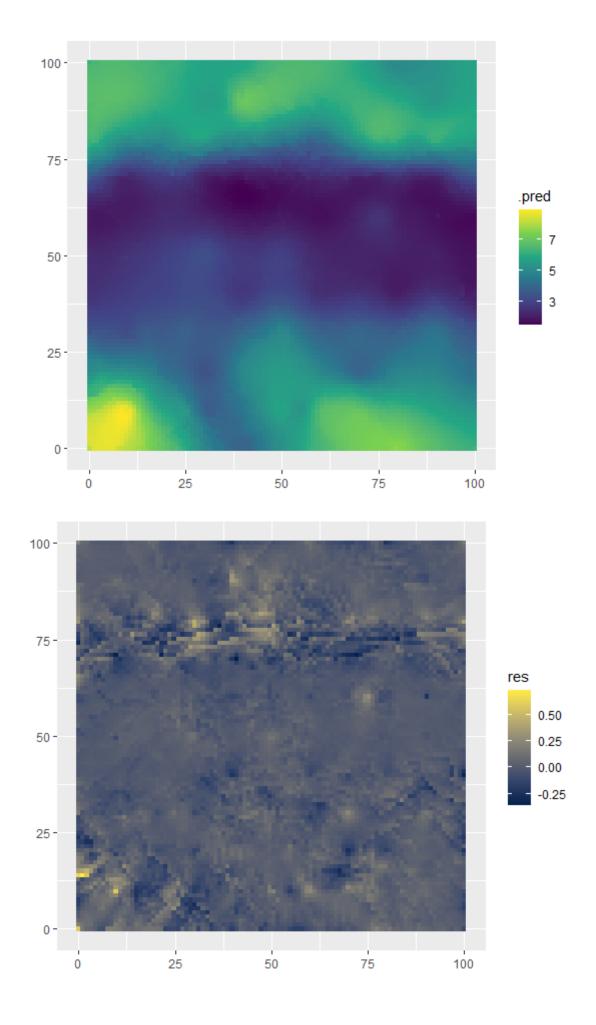


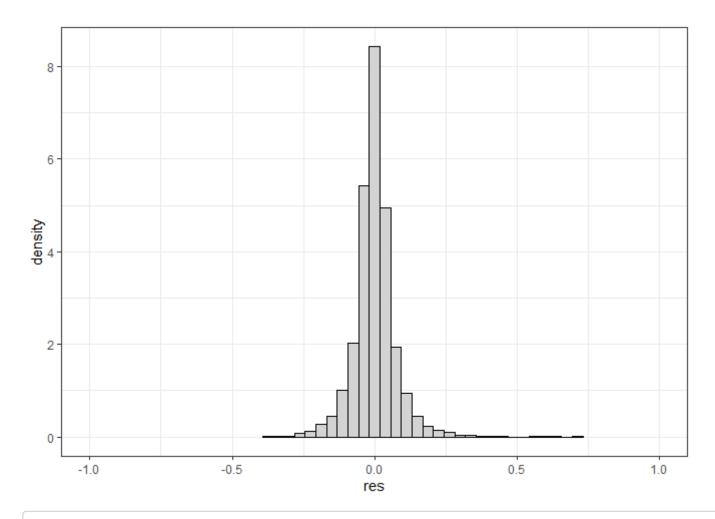


#> [1] "2017-03-15"

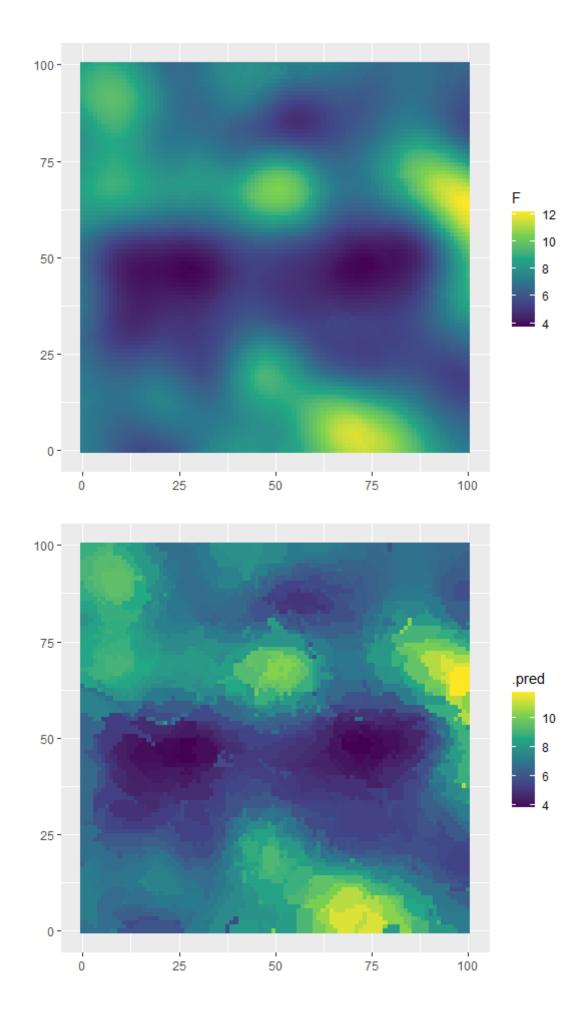


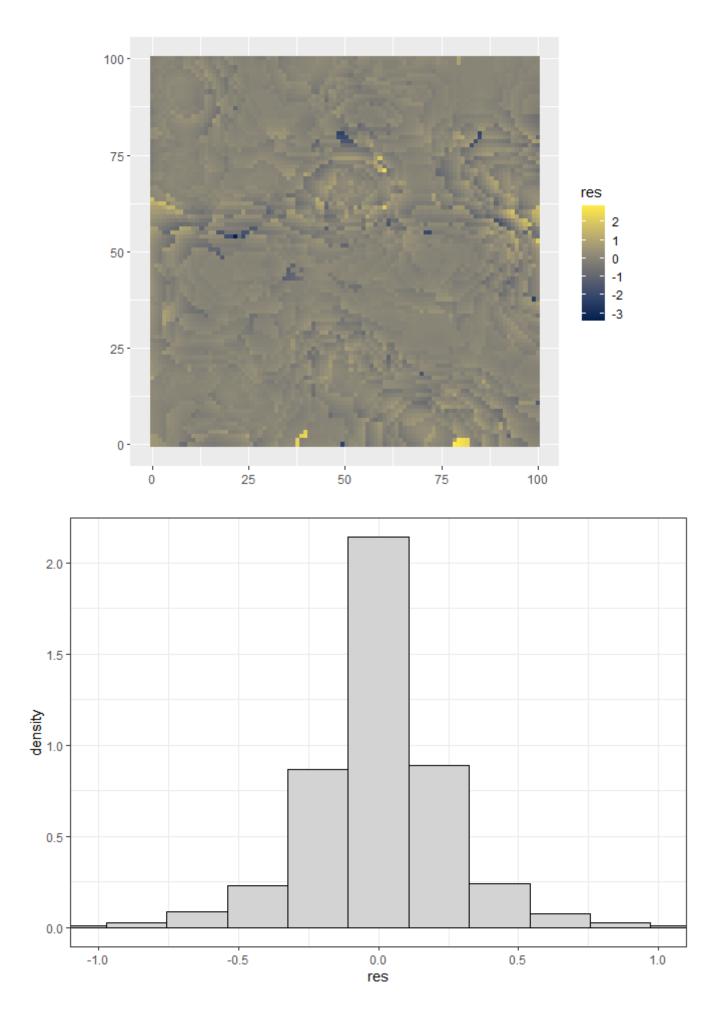


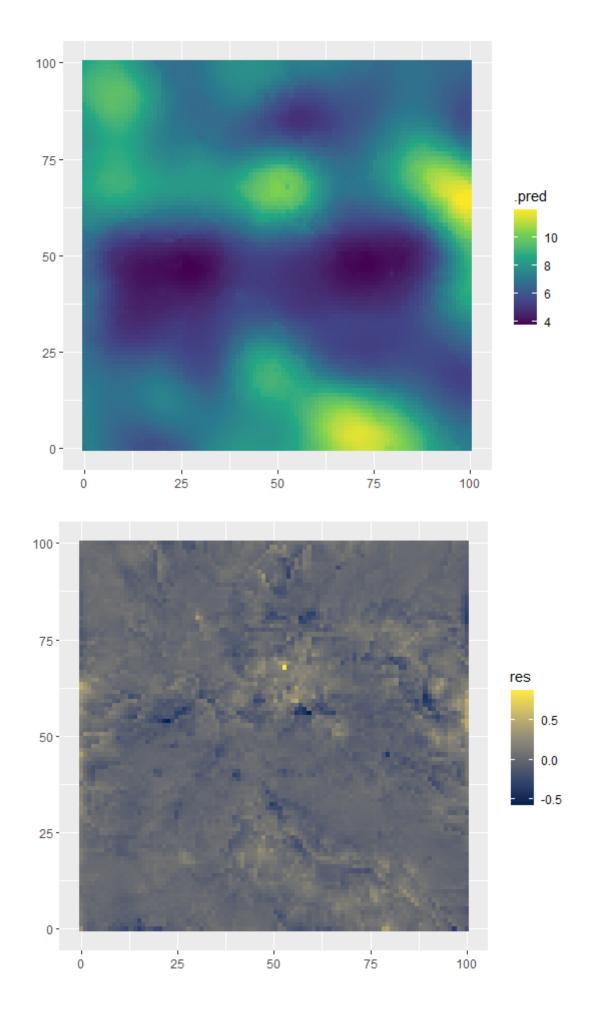


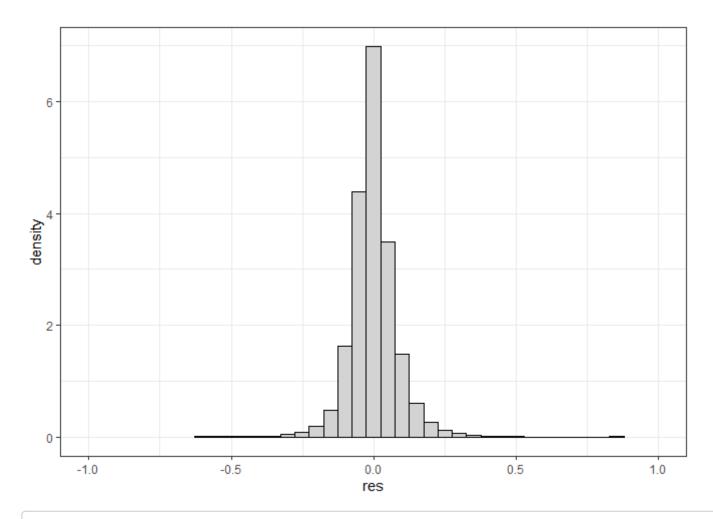


#> [1] "2017-02-17"

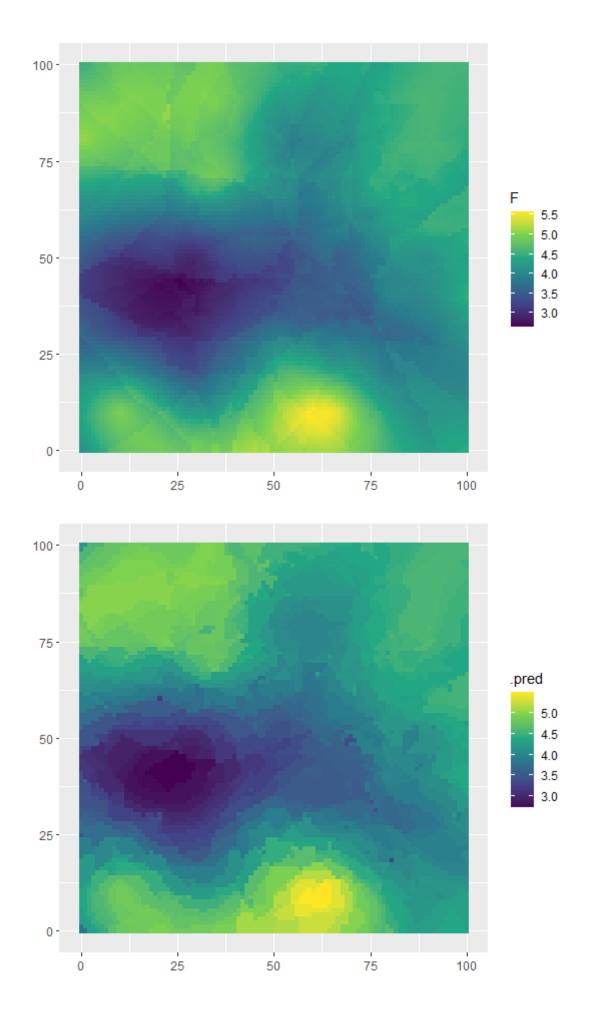


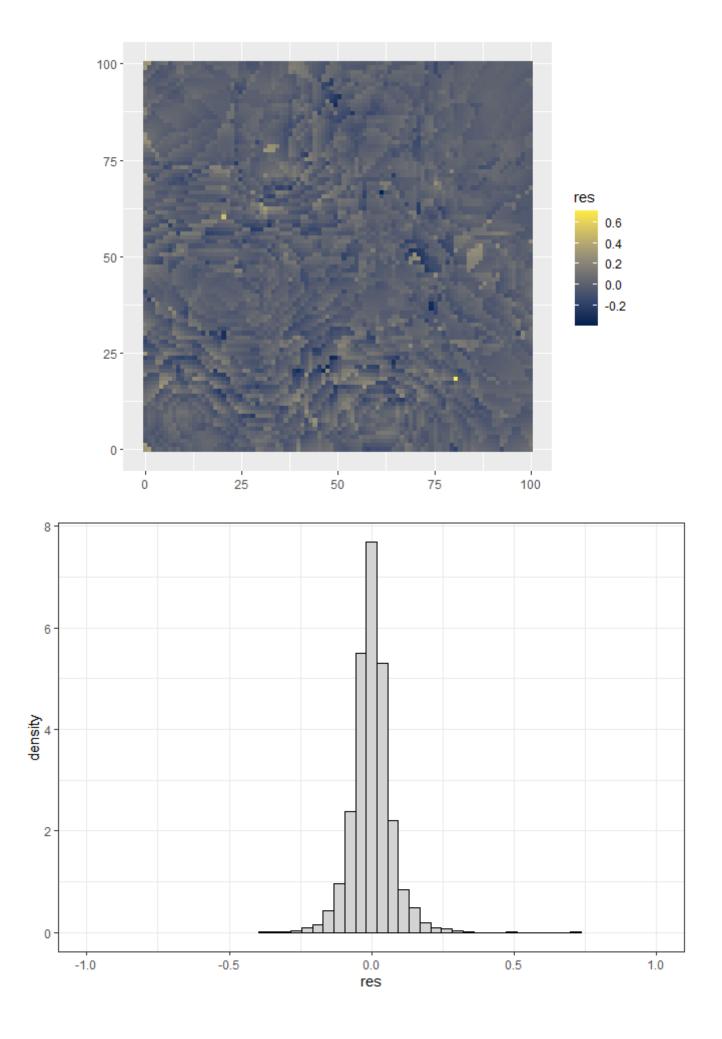


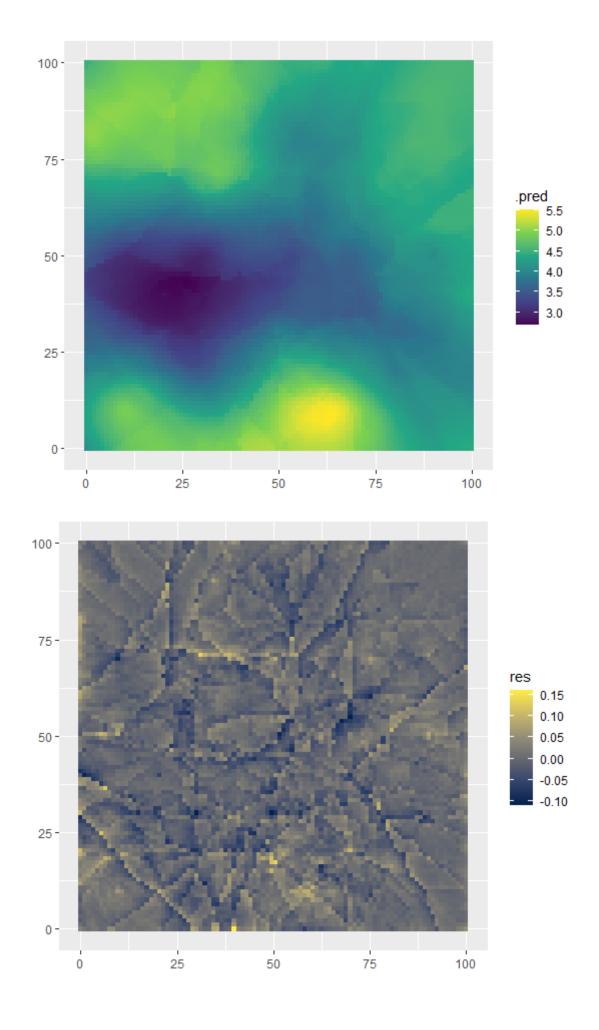


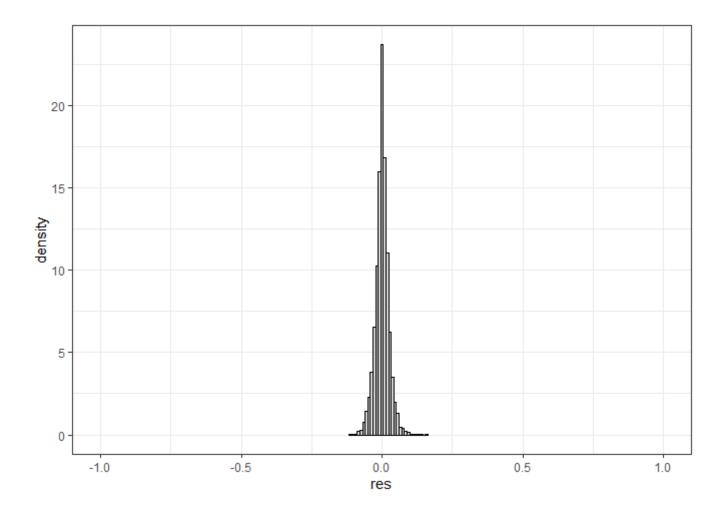


#> [1] "2017-06-17"





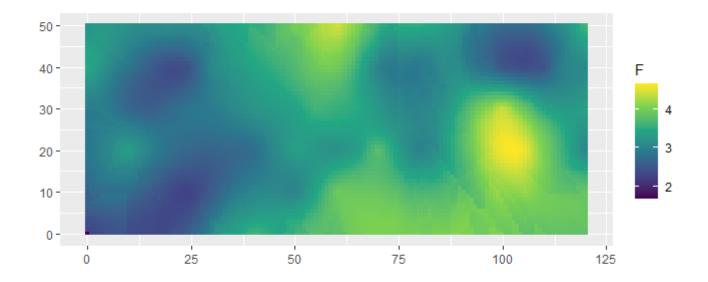


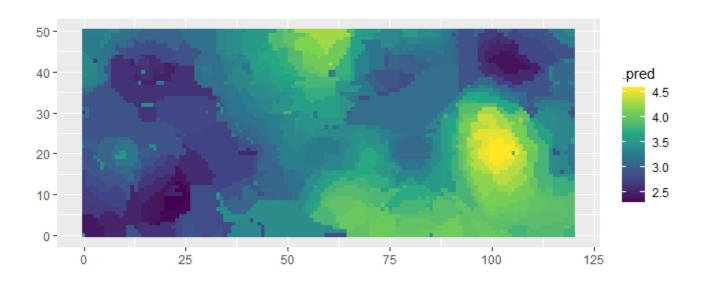


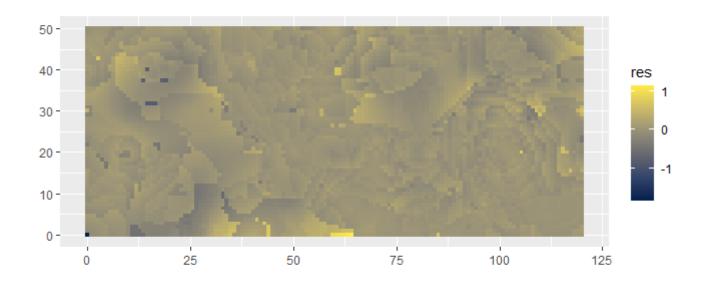
Criando os preditos para cada dia - SILVIPASTORIL

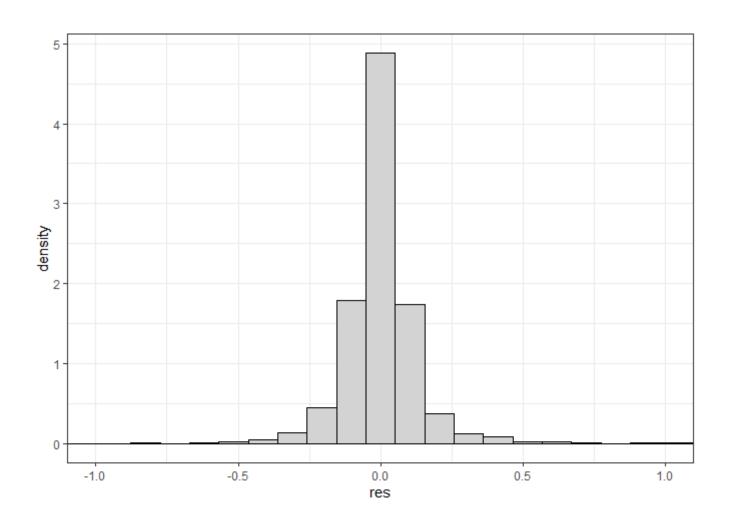
```
dias <- data sp$data %>% unique()
for(i in seq along(dias)){
 di <- dias[i]</pre>
 data sp %>% filter(data == di)
 file models sp <- list.files("models-3", pattern = "SP")</pre>
 file models sp <- grep(paste0(di), file models sp, value = TRUE)
 fco2 modelo load dt <- read rds(
   paste0("models-3/",grep("dt",file_models_sp,value=TRUE)
   ))
 fco2_modelo_load_rf <- read_rds(</pre>
   paste0("models-3/",grep("rf",file models sp,value=TRUE)
   ))
  # fco2 modelo load xgb <- read rds(
    paste0("models-3/",grep("xgb",file_models_sp,value=TRUE)
    ))
 yp dt <- predict(fco2 modelo load dt, new data = data sp %>%
                     filter(data == di))
 yp_rf <- predict(fco2_modelo_load_rf, new_data = data_sp %>%
                     filter(data == di))
  # yp_xgb <- predict(fco2_modelo_load_xgb, new_data = data_sp %>%
                  filter(data == di))
dis y < -50/55
dis x < -120/131
grid \leftarrow expand.grid(Y=seq(0,50,dis y), X=seq(0,120,dis x))
 mp krg <- tibble(grid, data sp %>%
                     filter(data == di)) %>%
   ggplot(aes(x=X,y=Y)) +
   geom tile(aes(fill = F)) +
   scale fill viridis c() +
   coord_equal()+labs(x="",y="")
 mp_dt <- tibble(grid, data sp %>%
                    filter(data == di), yp dt) %>%
   ggplot(aes(x=X,y=Y)) +
   geom tile(aes(fill = .pred)) +
   scale fill viridis c() +
   coord equal() +labs(x="",y="")
 mp rf <- tibble(grid, data sp %>%
                    filter(data == di), yp rf) %>%
   ggplot(aes(x=X,y=Y)) +
   geom tile(aes(fill = .pred)) +
   scale fill viridis c() +
   coord equal() +labs(x="",y="")
  mp dt res <- tibble(grid, data sp %>%
```

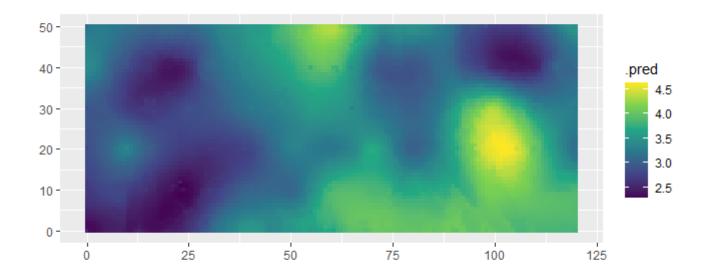
```
filter(data == di), yp dt) %>%
   mutate(res = F - .pred) %>%
   ggplot(aes(x=X,y=Y)) +
   geom tile(aes(fill = res)) +
   scale_fill_viridis_c(option = "E") +
   coord equal() +labs(x="",y="")
  hist_dt_res <- tibble(grid, data_sp %>%
                          filter(data == di), yp dt) %>%
   mutate(res = F - .pred) %>%
   ggplot(aes(x=res, y=..density..)) +
   geom histogram(color="black",fill="lightgray") +theme bw()+
   coord cartesian(xlim=c(-1,1))
 mp rf res <- tibble(grid, data sp %>%
                    filter(data == di), yp rf) %>%
   mutate(res = F - .pred) %>%
   qqplot(aes(x=X,y=Y)) +
   geom_tile(aes(fill = res)) +
   scale fill viridis c(option = "E") +
   coord_equal() +labs(x="",y="")
 hist_rf_res <- tibble(grid, data_sp %>%
                          filter(data == di), yp rf) %>%
   mutate(res = F - .pred) %>%
   ggplot(aes(x=res, y=..density..)) +
   geom histogram(color="black",fill="lightgray") +theme bw()+
   coord cartesian(xlim=c(-1,1))
 print(di)
 print(mp krg)
 print(mp dt)
 print(mp dt res)
 print(hist dt res)
 print(mp rf)
 print(mp rf res)
 print(hist rf res)
#> [1] "2017-02-03"
```

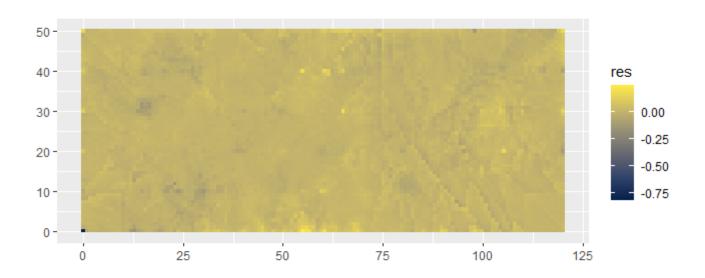


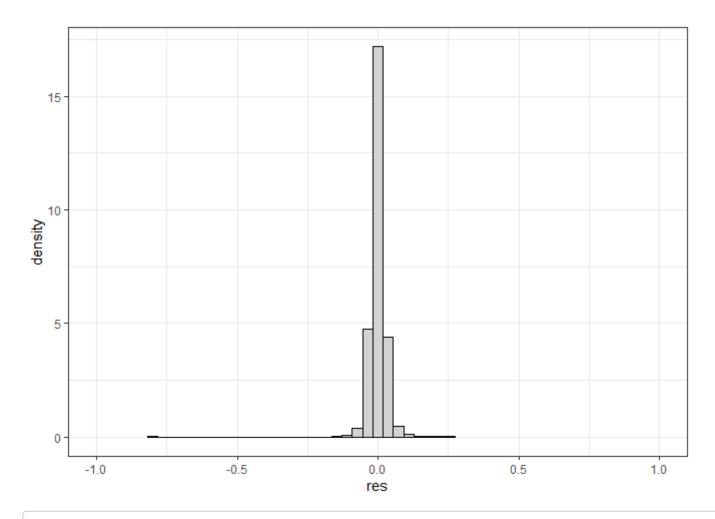




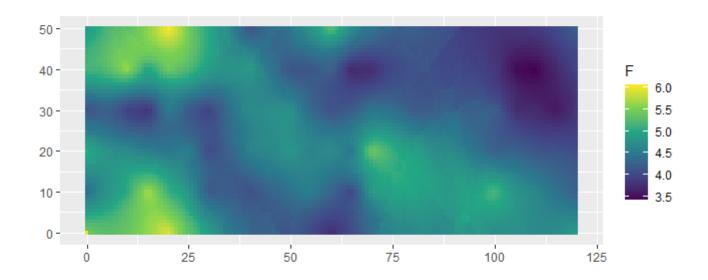


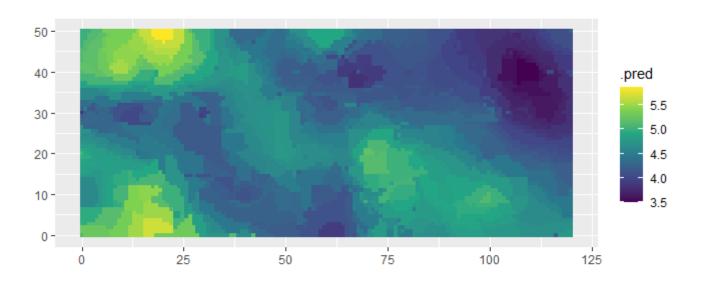


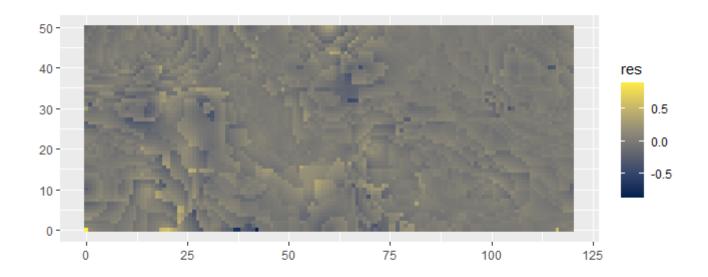


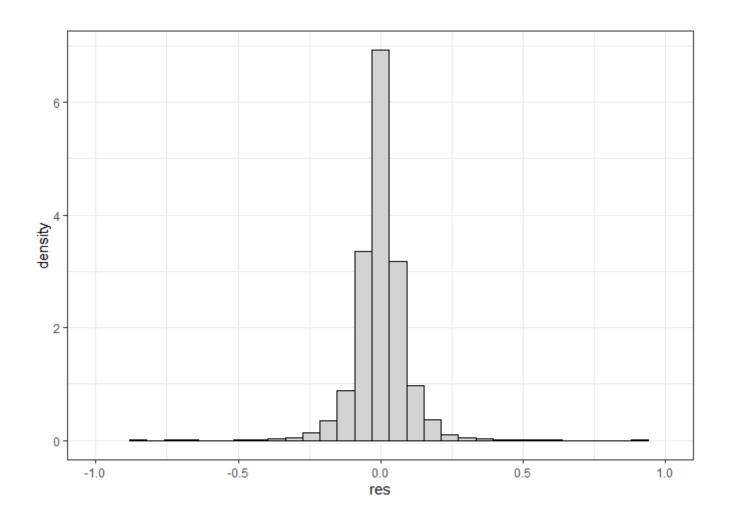


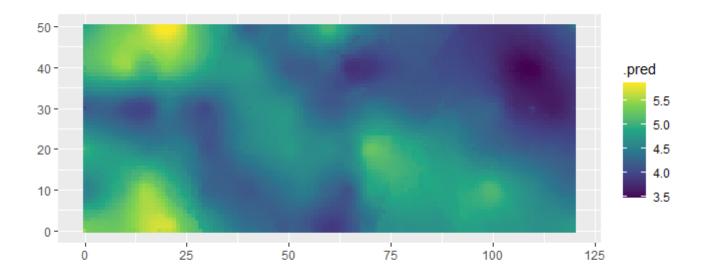
#> [1] "2017-03-03"

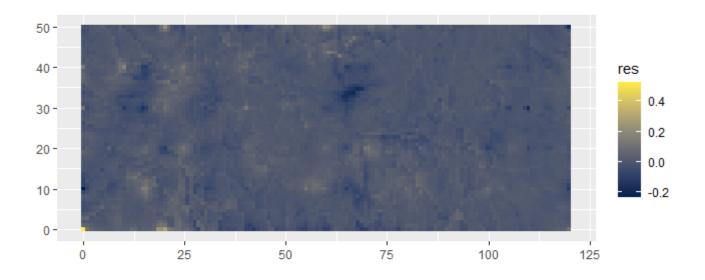


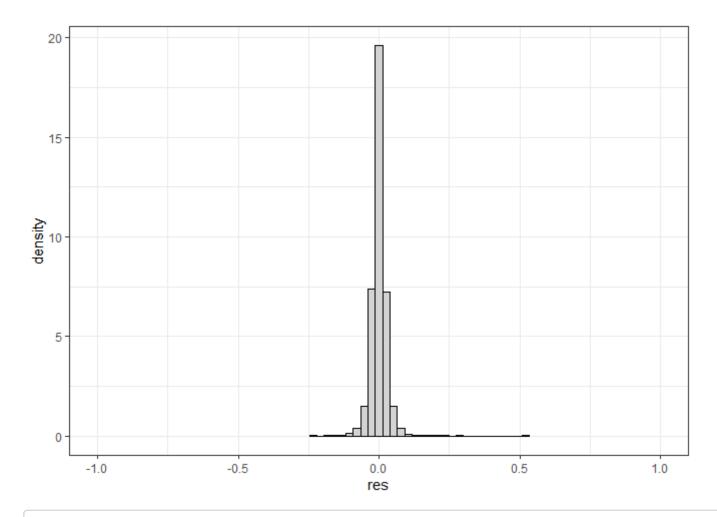




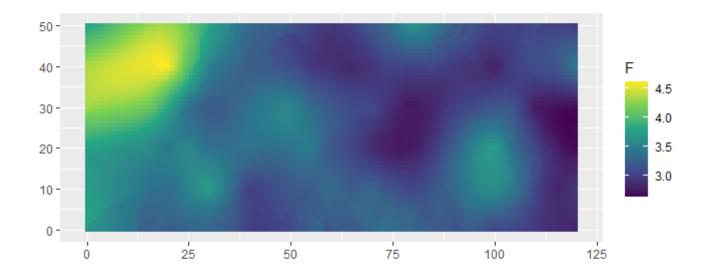


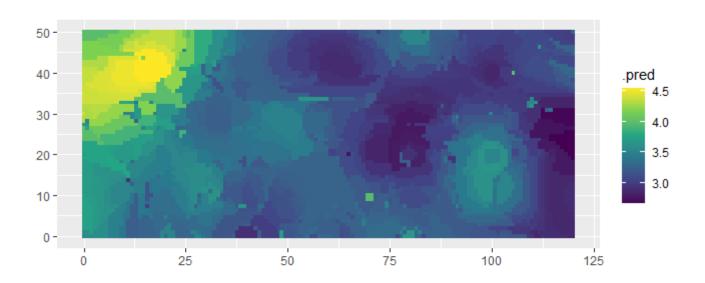


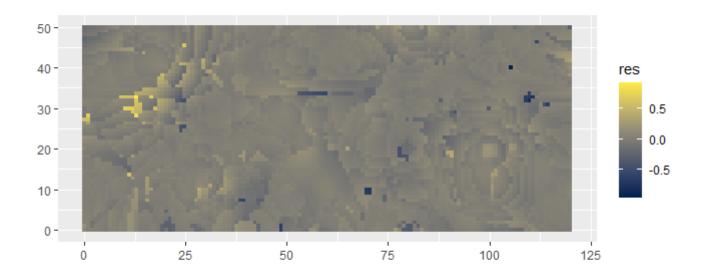


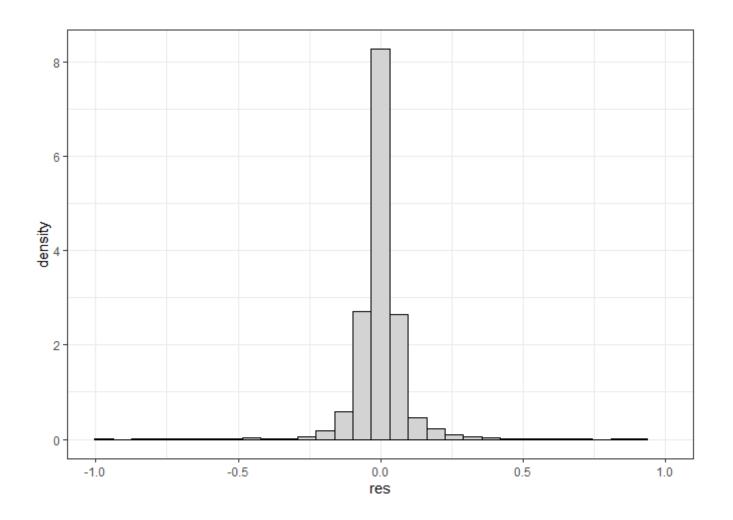


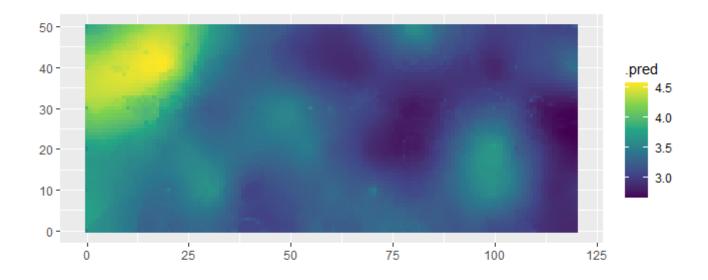
#> [1] "2017-06-03"

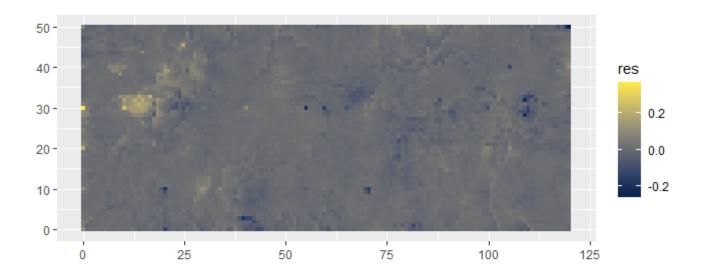


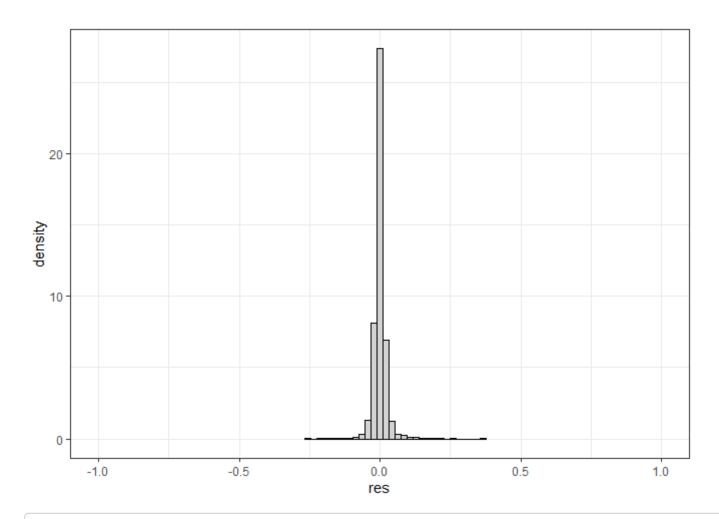




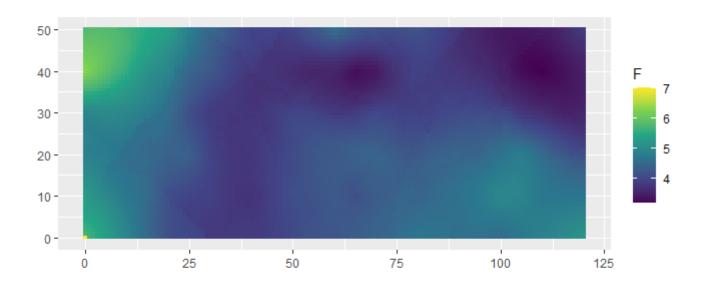


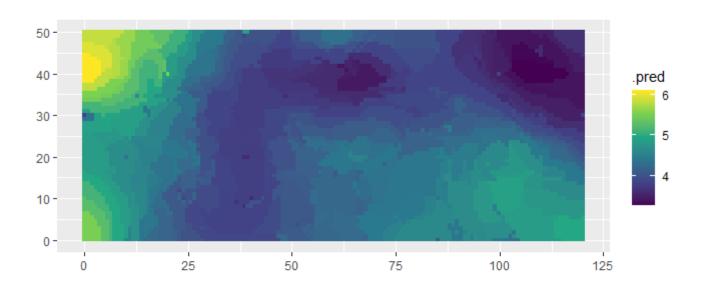


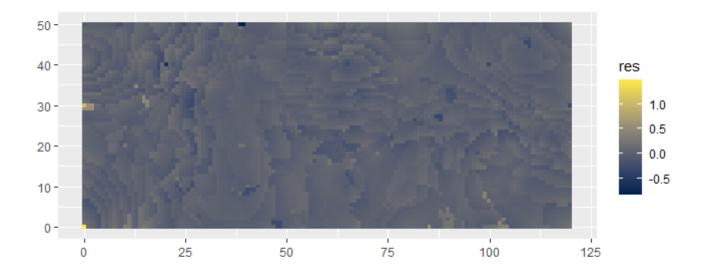


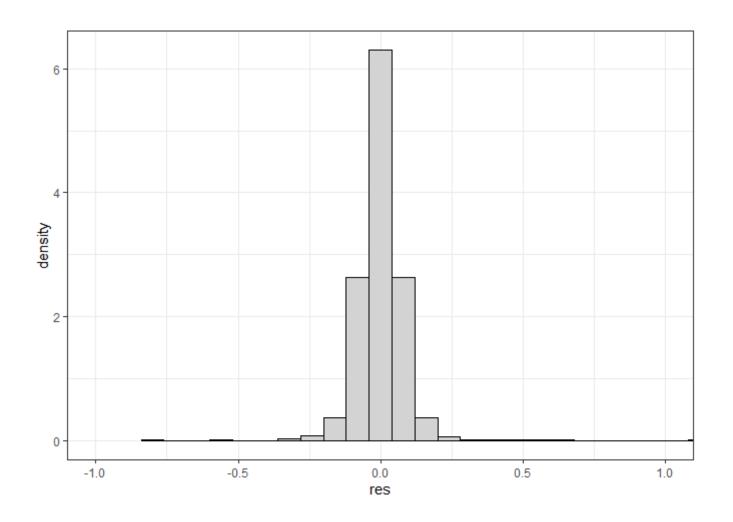


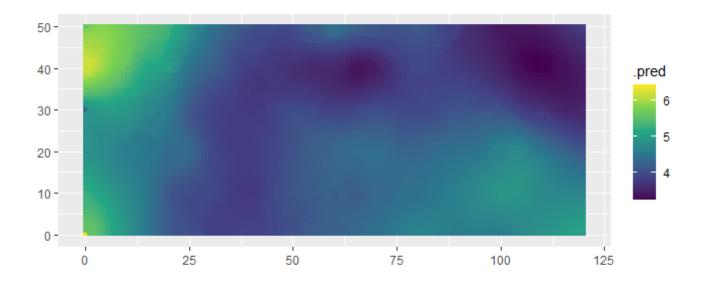
#> [1] "2017-03-08"

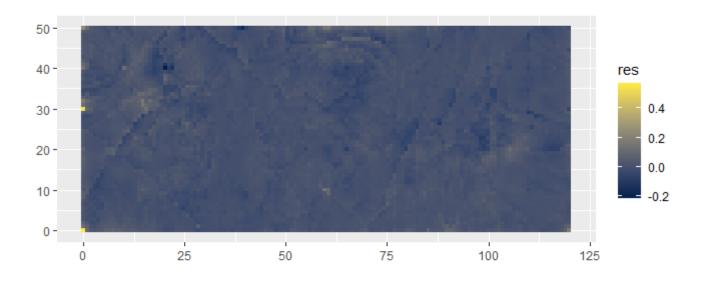


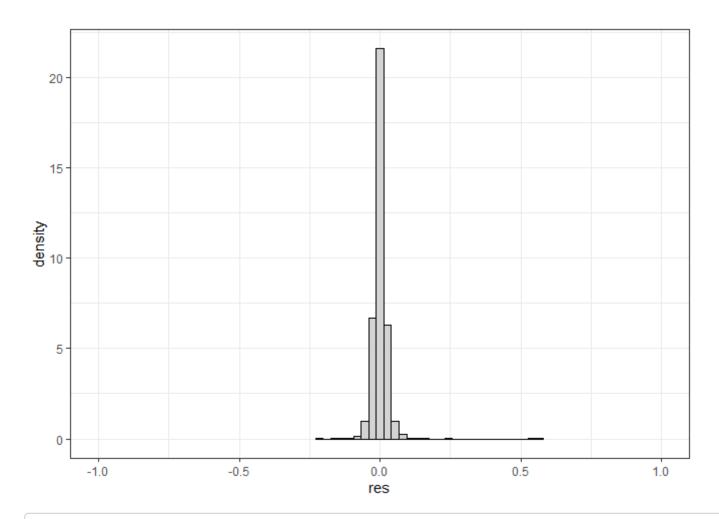




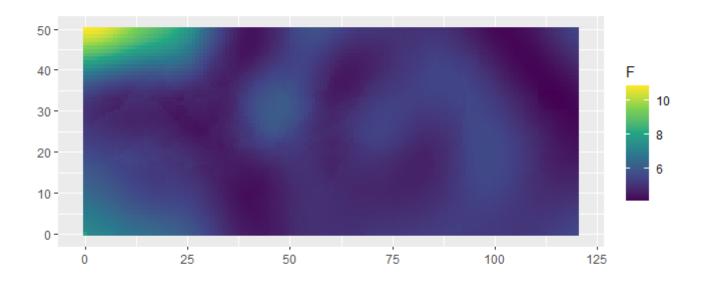


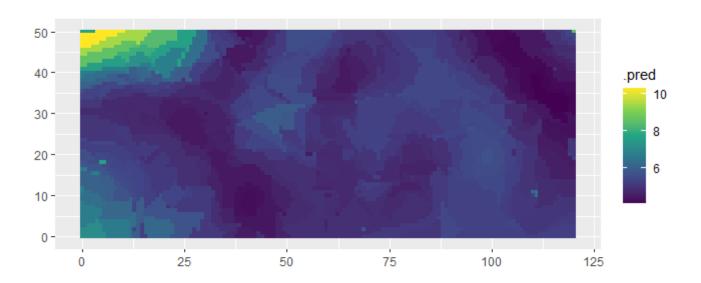


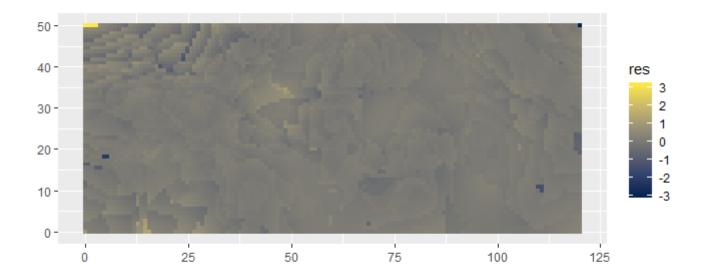


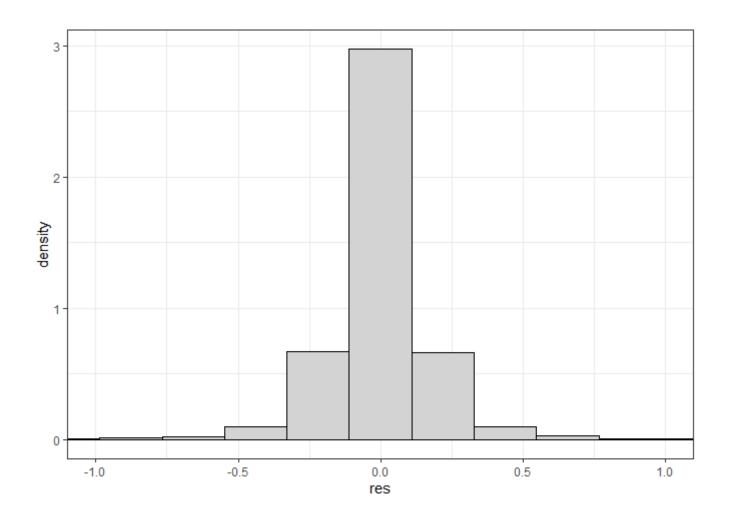


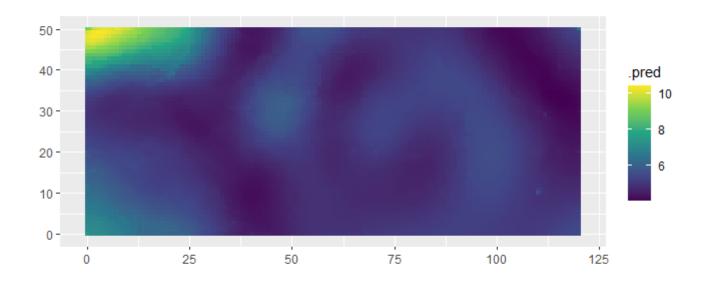
#> [1] "2017-02-09"

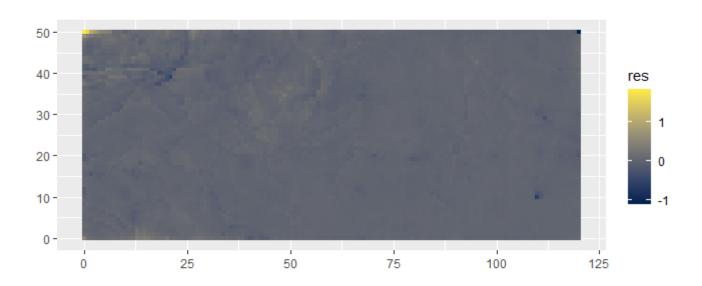


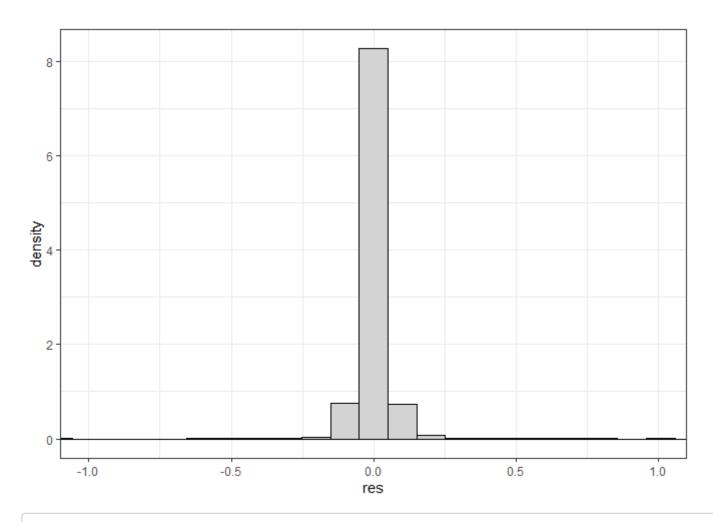




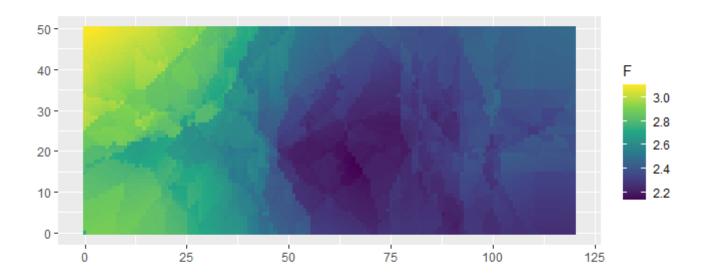


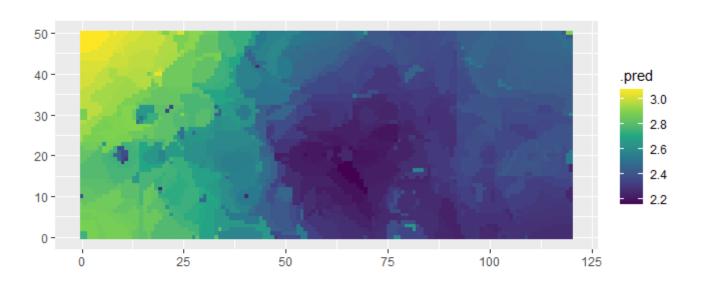


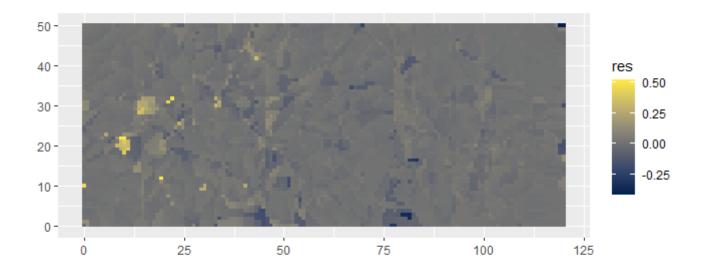


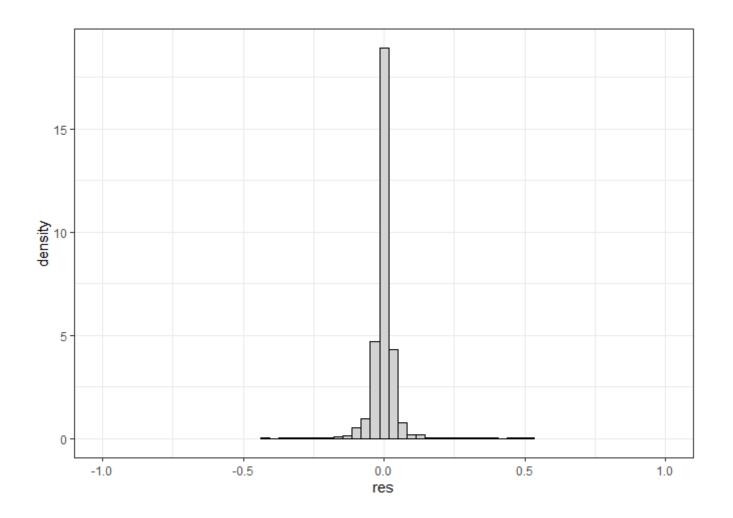


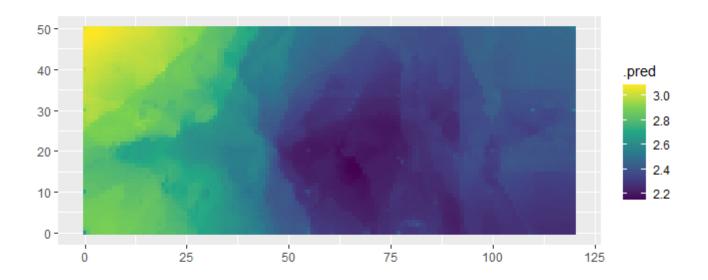
#> [1] "2017-06-10"

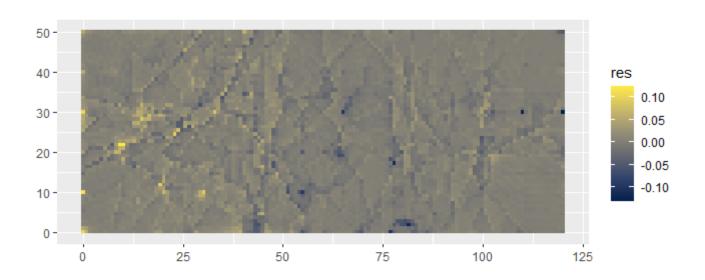


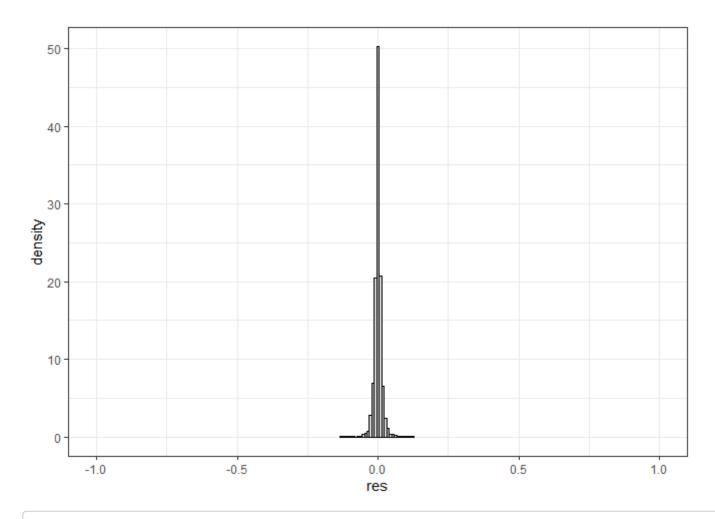




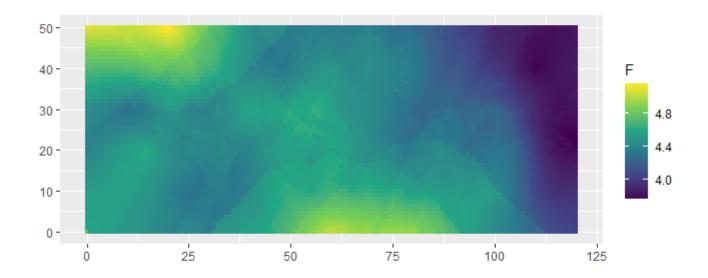


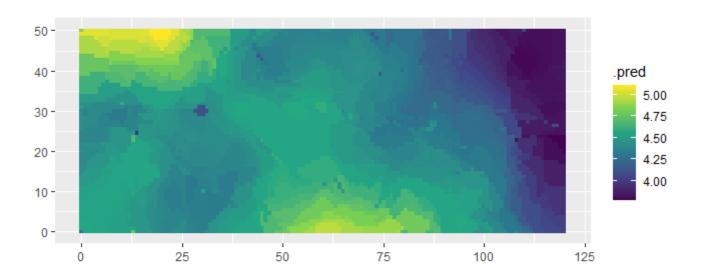


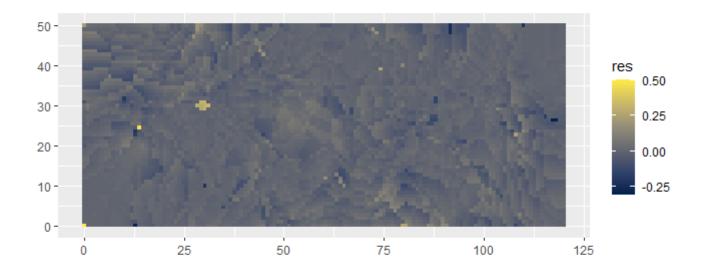


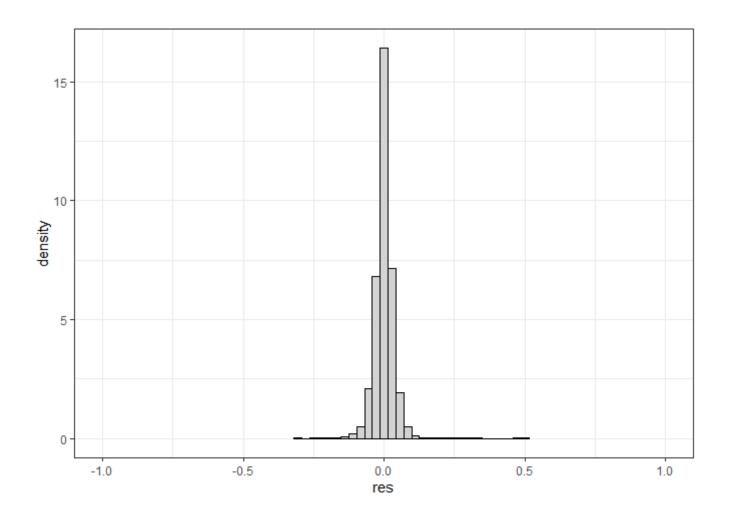


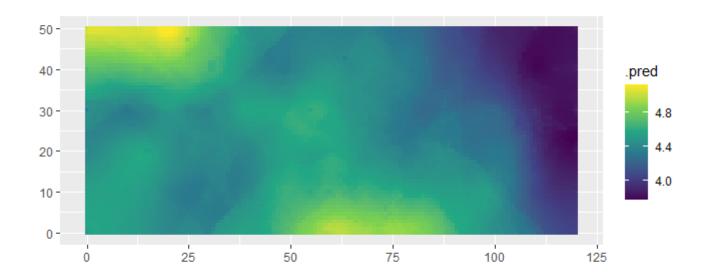
#> [1] "2017-03-17"

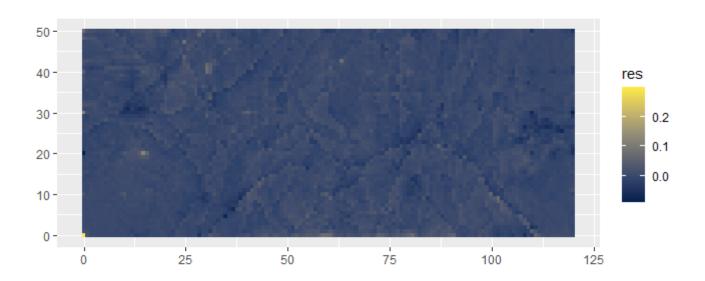


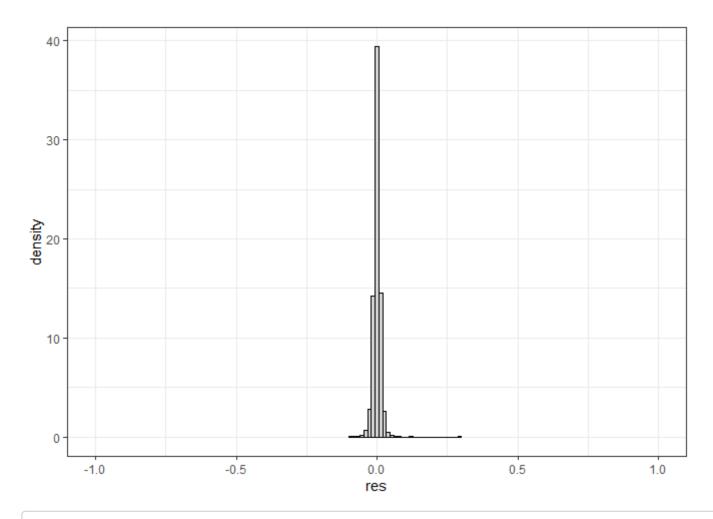




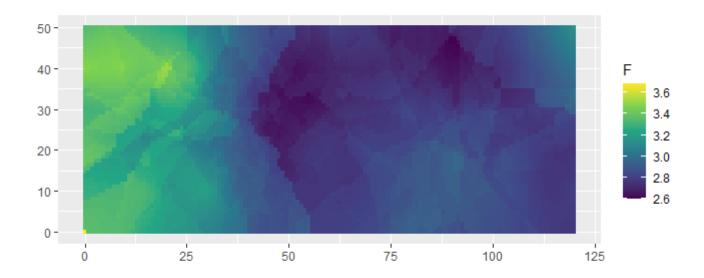


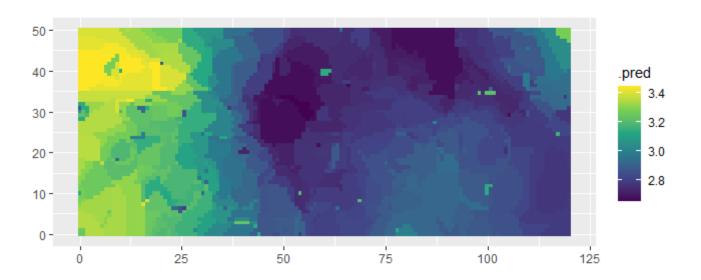


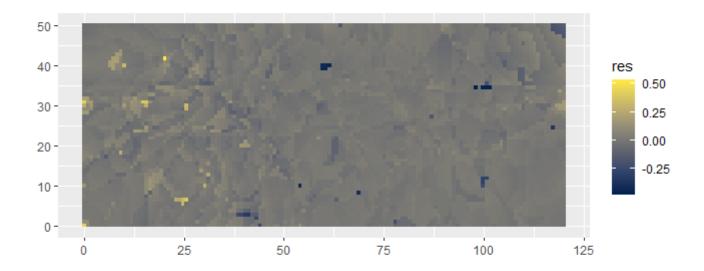


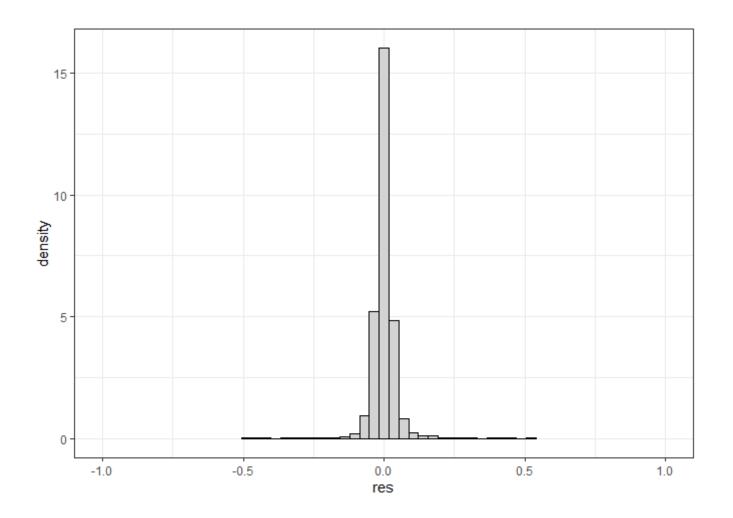


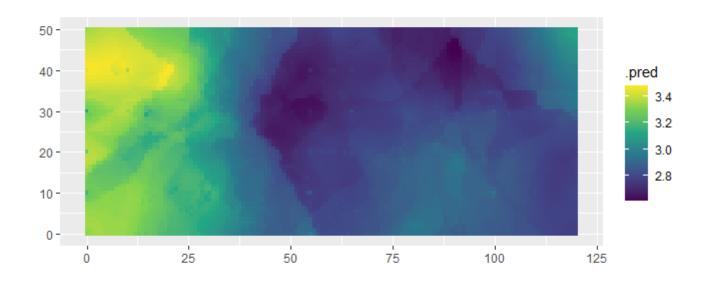
#> [1] "2017-06-17"

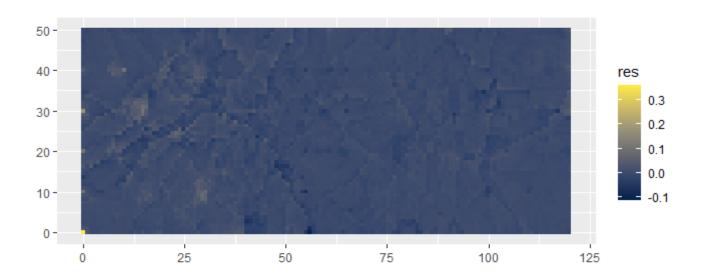


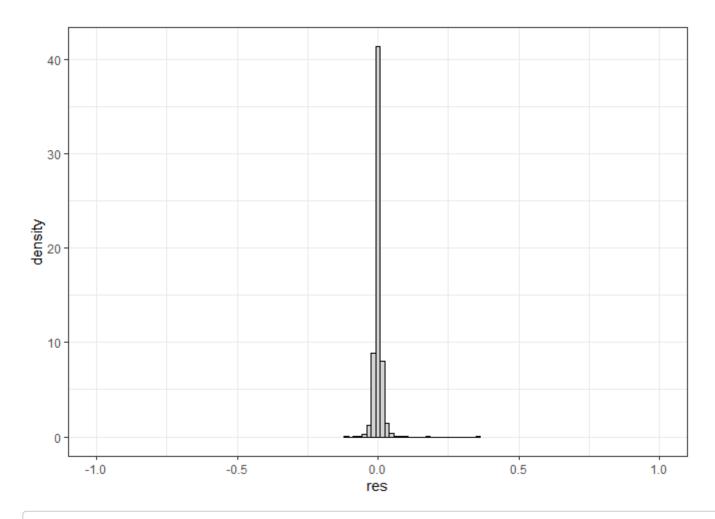




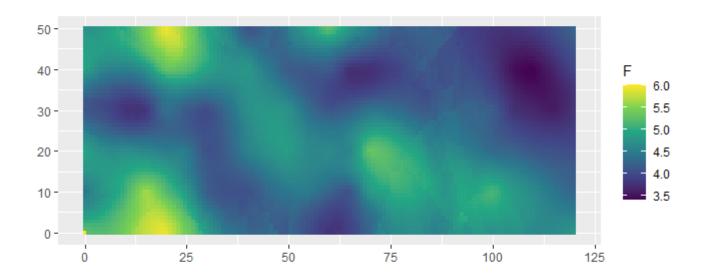


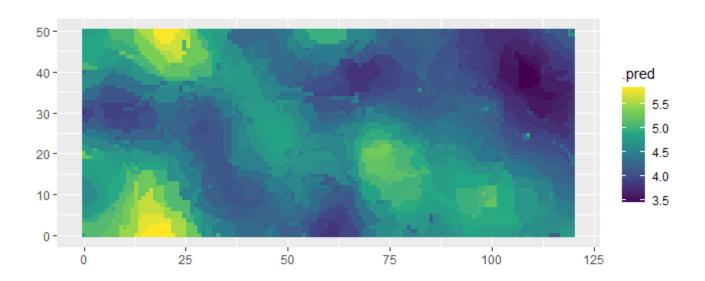


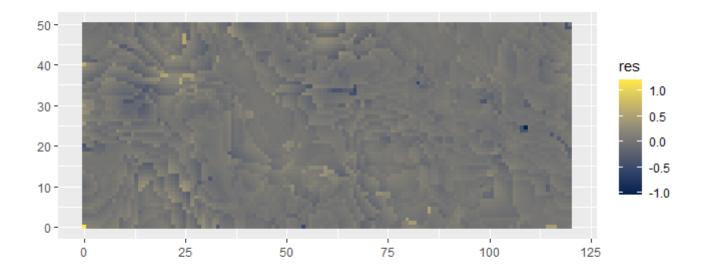


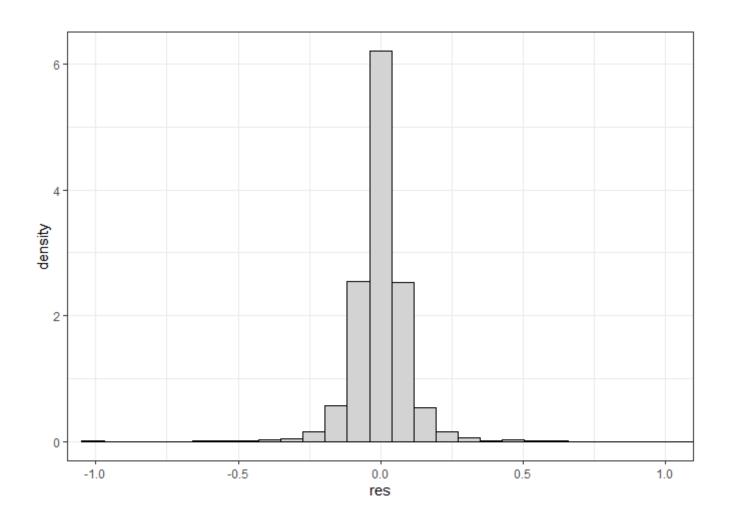


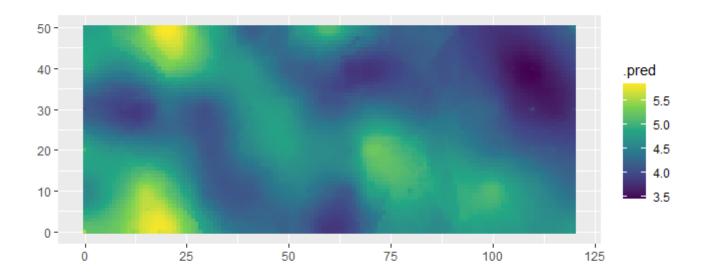
#> [1] "2017-02-22"

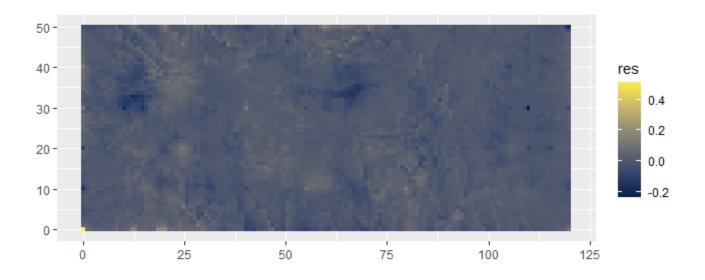


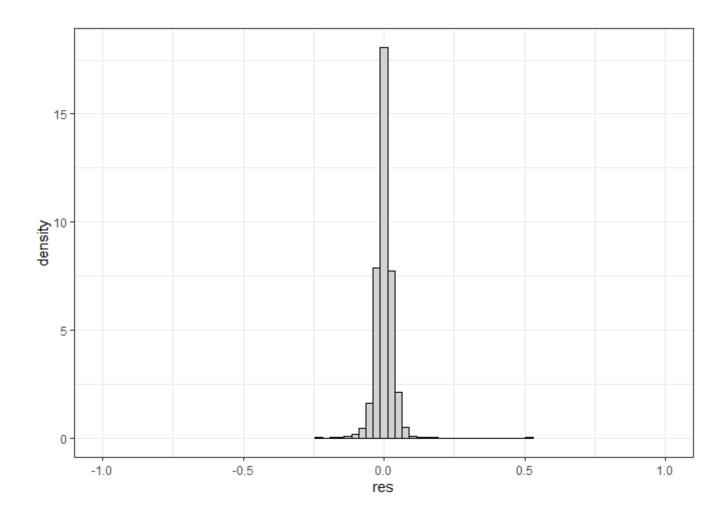












Aprendizado de Máquina

```
# # 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]
    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(
#
       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,
#
#
     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(
#
       min n = tune(),
      mtry = tune(),
#
#
       trees = tune()
#
     ) 응>응
#
       set mode("regression") %>%
#
       set engine("randomForest")
#
#
     fco2 rf wf <- workflow()</pre>
                                응>응
#
       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)
#
#
      fco2 test preds <- bind rows(
#
       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(\",\") \proonup \"))) +
#
        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$.pred)</pre>
#
     my r2 <- my r*my r
#
      my mse <- Metrics::mse(da$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
#
      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(
#
       mtry = 0.8,
#
        trees = tune(), # <-----
#
       min n = 5,
#
       tree depth = 4,
#
        loss reduction = 0, # lambda
#
       learn rate = tune(), # epsilon
#
       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)
#
#
      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 \times gb,
#
        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(
#
       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")
                                응>응
#
        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 \times gb \cdot 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(
#
       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)
#
```

```
#
      #### Grid
#
      fco2 xgb grid <- expand.grid(
        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(
#
       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$learn rate,
#
        sample size = tune()
#
      ) 응>응
#
        set mode("regression")
#
        set engine("xgboost", nthread = cores, counts = FALSE)
#
#
        fco2 xgb wf <- workflow() %>%
#
        add model(fco2 xgb model) %>%
#
        add recipe(fco2 recipe)
#
#
      fco2 xgb grid <- expand.grid(</pre>
        sample size = seq(0.5, 1.0, length.out = 2), ## <---
#
#
        mtry = seq(0.1, 1.0, length.out = 2) ## <---
#
      )
#
#
      fco2 xgb tune grid <- fco2 xgb wf
#
        tune grid(
#
          resamples = fco2 resamples,
#
          grid = fco2 xgb grid,
          control = control_grid(save_pred = TRUE, verbose = FALSE, allow par = TRUE),
#
#
          metrics = metric set(rmse)
#
#
      fco2 xgb select best passo4 <- fco2 xgb tune grid %>% select best(metric = "rm
se")
#
#
      # passo 5
#
      fco2 xgb model <- boost tree(
#
       mtry = fco2 xgb select best passo4$mtry,
#
        trees = tune(),
#
        min n = fco2 xgb select best passo2$min n,
#
        tree depth = fco2 xgb select best passo2$tree depth,
```

```
loss reduction = fco2 xgb select best passo3$loss reduction,
#
        learn rate = tune(),
#
        sample size = fco2 xgb select best passo4$sample size
#
      ) 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>
#
       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 \times gb \cdot 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 = "rm
se")
#
#
      ## 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)</pre>
```

```
#
      fco2 test preds <- bind rows(
#
        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(\", \") \sim ~"))) +
        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
#
      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$F,
                                 da$.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,
                              RMSE=my rmse, MAE=my mae, MAPE=my mape)
      print(data.frame(vector of metrics))
#
# }
```

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])
# }
```