

# Evaluating the multiple forecasters

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## Data pre-loading and processing

```
library(here)
library(covidcast)
library(epiprocess)
library(zoltr)
library(tidyverse)
library(ggsci)
library(tsibble)
library(covidHubUtils)
library(lubridate)
library(rlang)
library(patchwork)
library(pROC)
here::i_am(path = "notebooks/multiple-forecaster.Rmd")
source(here("R", "utils.R"))
# (settings <- get_settings(start_date = "2020-06-01", end_date = "2022-03-01"))
theme_set(theme_bw())
```

First let's set some parameters and constants.

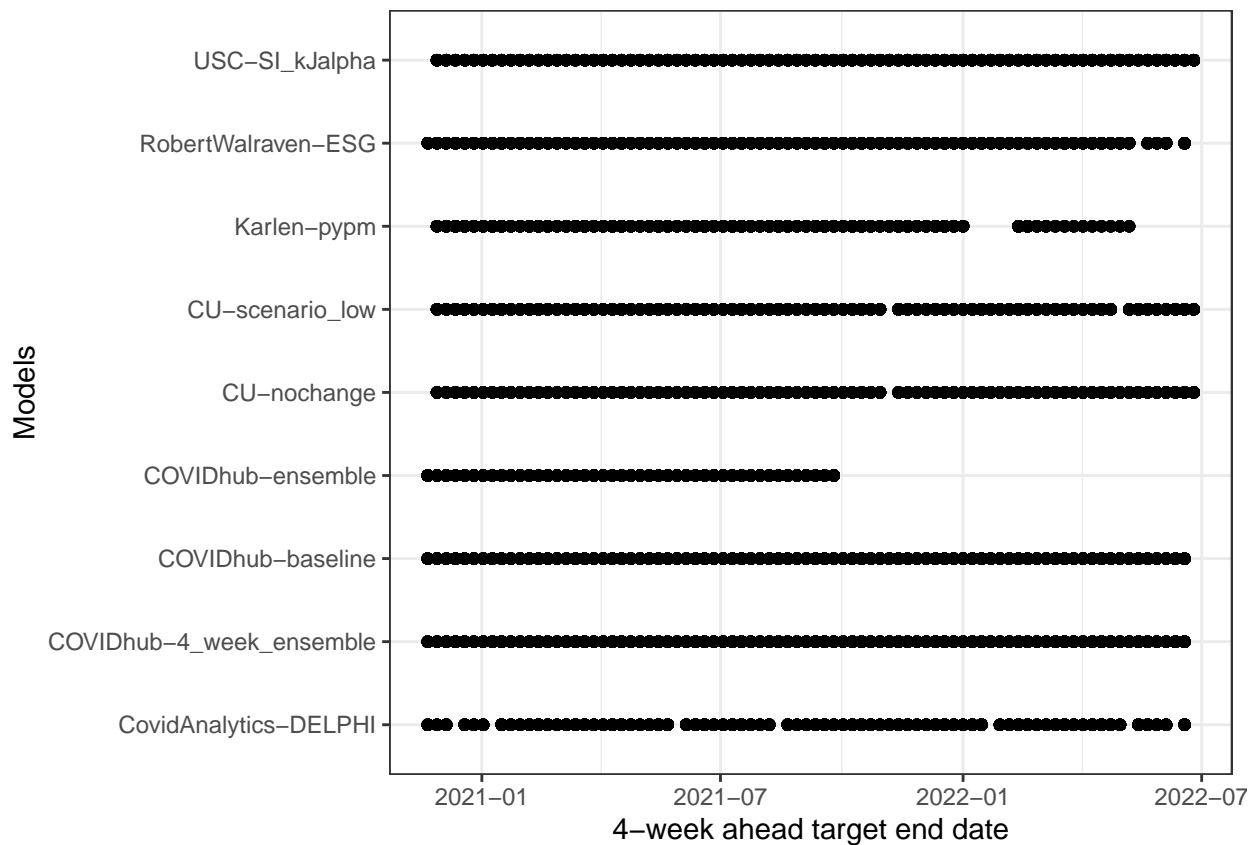
```
d_range <- seq(ymd("2020-11-01"), ymd("2022-06-01"), by = 7)
curr_date <- "2022-06-01"
h <- 4
wk_ahead <- 1
inc_case_targets <- paste(1:h, "wk ahead inc case")
surge_thresh <- 0.5
min_inc <- 20
```

## Loading forecasts and underlying data

Using `covidHubUtils` and `zoltar` we load underlying data as well as forecasts for models available in the US hub. Here we retrieve US states incident case forecasts 1-4 weeks in advance.

```
models <- c("COVIDhub-baseline", "RobertWalraven-ESG", "CovidAnalytics-DELPHI", "USC-SI_kJalpha",
            "COVIDhub-4_week_ensemble", "CU-nochange", "Karlen-pypm", "CU-scenario_low",
            "COVIDhub-ensemble")
if (!file.exists(here("data", "pred_data_20220608.rds"))){
  pred_case <- load_forecasts(
    models = models,
    dates = d_range,
    date_window_size = 6,
    locations = state.name,
    types = "point",
    targets = inc_case_targets,
    source = "zoltar",
    verbose = FALSE,
    as_of = curr_date,
    hub = c("US")
  )
  saveRDS(pred_case, file = here("data", "pred_data_20220608.rds"))
} else {
  pred_case <- readRDS(file = here("data", "pred_data_20220608.rds"))
}

ggplot(pred_case %>% filter(horizon == 4), aes(x = model, y = target_end_date)) +
  geom_point() +
  labs(x = "Models", y = "4-week ahead target end date") +
  coord_flip()
```



Let's load the corresponding truth data

```
truth_data <- load_truth(
  truth_source = "JHU",
  target_variable = "inc case",
  locations = state.name
)

true_range <- pred_case %>% pull(target_end_date) %>%
  unique() %>%
  lubridate::as_date()

# convert truth_data to epi_df
truth_epidf <- truth_data %>%
  select(-c(model, location, target_variable, location_name,
            abbreviation, full_location_name)) %>%
  dplyr::rename("time_value" = "target_end_date") %>%
  filter(time_value %in% true_range) %>%
  as_epidf(geo_type = "state")

truth_epidf

## An `epi_df` object, with metadata:
## * geo_type = state
## * time_type = day
## * as_of = 2022-06-28 14:41:35
##
## # A tibble: 4,350 x 5
```

```
##   geo_value time_value value population geo_type
## * <chr>      <date>      <dbl>      <dbl> <chr>
## 1 al        2020-10-31  9782      4903185 state
## 2 al        2020-11-07  9905      4903185 state
## 3 al        2020-11-14 12325      4903185 state
## 4 al        2020-11-21 14865      4903185 state
## 5 al        2020-11-28 14285      4903185 state
## 6 al        2020-12-05 22596      4903185 state
## 7 al        2020-12-12 25252      4903185 state
## 8 al        2020-12-19 27063      4903185 state
## 9 al        2020-12-26 23554      4903185 state
## 10 al       2021-01-02 26000      4903185 state
## # ... with 4,340 more rows
```

## Surge classification using relative change growth rate formulation

We define surge for a given date using relative change growth rate formulation times the bandwidth using the implementation from `epiprocess`:

$$\frac{1}{h} * \left( \frac{\bar{B}}{\bar{A}} - 1 \right) = \frac{1}{h} * \left( \frac{\bar{B} - \bar{A}}{\bar{A}} \right) = \frac{1}{h} * \left( \frac{(h)^{-1} \left( \sum_{t=T+1}^{T+h} Y_t - \sum_{t=T+1-h}^T Y_t \right)}{(h)^{-1} \sum_{t=T+1-h}^T Y_t} \right) = \frac{1}{h} R_{T+h}^h$$

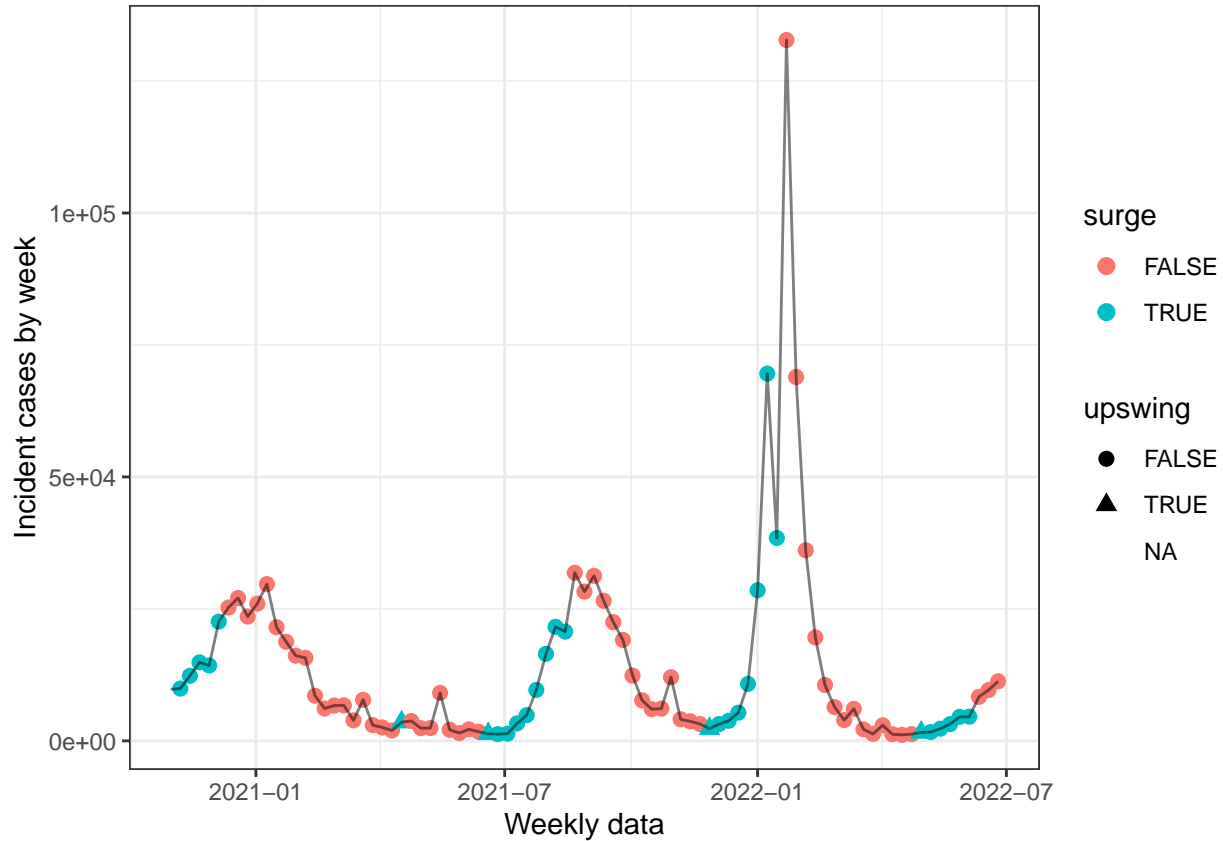
A surge is defined for time-point  $T$  as the difference in cumulative incident cases between the periods of  $T + 1$  and  $T + h$  and  $T$  and  $T - h$ . As such, an  $h$ -week ahead forecaster is a nowcaster of whether or not we're currently in a surge. Here, we classify for each week whether it is a surge based on it's  $h$ -week-ahead and  $h$ -week-prior data. An upswing is defined as the point at the beginning of a surge. Here we used the threshold of 50% increase in cumulative difference in the  $h$ -week-ahead compared to  $h$ -week-prior data.

```
truth_epidf <- truth_epidf %>%
  mutate(gr = growth_rate(y = value, method = "rel_change", h = h) * h) %>%
  mutate(surge = case_when(
    gr >= surge_thresh & value >= min_inc ~ TRUE,
    TRUE ~ FALSE
  ))

truth_epidf <- truth_epidf %>%
  group_by(geo_value) %>% epi_slide(~{
    bef <- .x$surge[1]
    focal <- .x$surge[2]
    if (is.na(focal)){
      out <- NA
    } else {
      if (bef == FALSE & focal == TRUE){
        out <- TRUE
      } else if (bef == TRUE & focal == FALSE){
        out <- FALSE
      } else {
        out <- FALSE
      }
    }
    return(out)
  }, n = 2 * 7 * 1, align = "right", new_col_name = "upswing")
```

```
ggplot(truth_epidf %>% filter(geo_value == "al"), aes(x = time_value, y = value)) +
  geom_point(aes(col = surge, shape = upswing), size = 2.5) + geom_line(alpha = 0.5) +
  labs(x = "Weekly data", y = "Incident cases by week")
```

## Warning: Removed 1 rows containing missing values (geom\_point).



## Nowcasting surges

Here we use the following procedures for a focal timepoint  $T$  and bandwidth  $h$  (for example, 4)

1. We take a time period from  $T + 1 - h$  to  $T + h$ . For example at 2021-01-23, we'd be taking the period from 2021-01-02 to 2021-02-20.
2. We take the truth period to be from  $T + 1 - h$  to  $T$  (inclusive) and the forecasting period to be from  $T + 1$  to  $T + h$ . The truth period would have real underlying incident cases while the forecasting period has forecast incident cases at times  $1-h$  weeks ahead. Due to the forecasting date being on Monday instead of exactly one week before the proposed target date, we take forecast incident values from the forecast date closest to the time period defined at  $T + 1$ . For example, at 2021-01-23, we would take forecast values for 2021-01-30 onwards from a forecast date of 2021-01-25.
3. We compute the growth rate at time  $T$  using these two periods as per the formula above
4. We then classify periods as surges using the definition and thresholds defined above.

```
forecast <- pred_case %>% dplyr::rename("pred" = value, "time_value" = target_end_date)
actual <- truth_epidf %>% dplyr::rename("obs" = value)
```

```
combined <- left_join(actual, forecast) %>%
  select(geo_value, population, geo_type, obs, pred,
         time_value, forecast_date, model, horizon, surge, upswing)
```

```
## Joining, by = c("geo_value", "time_value", "population", "geo_type")
```

```
combined <- combined %>% ungroup() %>% as_epi_df()
head(combined)
```

```
## # A tibble: 6 x 11
##   geo_value time_value population geo_type   obs   pred forecast_date model
##   <chr>      <date>      <dbl> <chr>    <dbl> <dbl> <date>      <chr>
## 1 ak        2020-10-31      731545 state    2812 1725  2020-10-26 COVIDhub-b~
## 2 ak        2020-10-31      731545 state    2812 1557. 2020-10-26 RobertWalr~
## 3 ak        2020-10-31      731545 state    2812 1039  2020-10-26 CovidAnaly~
## 4 ak        2020-10-31      731545 state    2812 1664  2020-10-26 COVIDhub-4~
## 5 ak        2020-10-31      731545 state    2812 1664  2020-10-26 COVIDhub-e~
## 6 ak        2020-11-07      731545 state    2951 1725  2020-10-26 COVIDhub-b~
## # ... with 3 more variables: horizon <chr>, surge <lgl>, upswing <lgl>
```

Let's define the slide function

```
# this function combines real case counts from time points t-h to t and forecasted
# case counts from t+1 to t+h. Growth rate at time t is then estimated using the relative change
# method
mismatch_slide <- function(slide_df, h){
  query_dates <- slide_df %>% pull(time_value) %>% unique()
  req_len <- h * 2
  # if not enough weeks for prediction
  if (length(query_dates) != req_len){
    # this is an exception for when the dates are truncated at the end and beginning of the interval
    out <- NA_real_
  } else {
    # true dates, pred dates and ref dates
    t_date <- query_dates[1:(req_len - h)]
    ref <- tail(t_date, n = 1)
    p_date <- query_dates[(req_len - h + 1):req_len]
    #print(query_dates)
    #print(ref)
    f_date <- slide_df %>%
      filter(time_value == head(p_date, n = 1) & horizon == 1) %>%
      pull(forecast_date) %>% unique()

    if (length(f_date) == 0){
      # exceptions where there are no relevant forecasts for when the time step 1 ahead of
      # the reference date
      out <- NA_real_
    } else {
      # get predicted values
      pred <- slide_df %>% filter(forecast_date == f_date) %>%
        dplyr::pull(pred)

      if (length(pred) < h){
        # this exception is when the model does not forecast $h$ weeks in advance
        # (e.g. h = 4 but horizon only extends to 3)

```

```

    out <- NA_real_
  } else {
    # get true values
    obs <- slide_df %>% filter(time_value %in% t_date) %>%
      select(time_value, obs) %>%
      distinct() %>% pull(obs)

    # put everything in an epi_df for posterity
    new_df <- tibble(
      time_value = c(t_date, p_date),
      value = c(obs, pred),
      geo_value = "placeholder"
    ) %>% as_epi_df()

    out <- new_df %>%
      mutate(gr_pred = growth_rate(y = value, h = h, method = "rel_change") * h) %>%
      filter(time_value == ref) %>% pull(gr_pred)
  }
}
return(out)
}

```

Let's use this with `epi_slide` function and loop through all the models

```

if (!file.exists(here("output", "calc_gr.rds"))){
  # this step takes about 20 minutes
  begin <- Sys.time()
  gradient_calc <- combined %>%
    group_by(geo_value, model) %>%
    epi_slide(~mismatch_slide(.x, h = h), n = 2 * 7 * h, align = "center", new_col_name = "pred_gr")
  dplyr::ungroup() %>%
  dplyr::select(-c(forecast_date, horizon, pred)) %>% distinct()
  end <- Sys.time()
  print(end - begin)
  saveRDS(gradient_calc, file = here("output", "calc_gr.rds"))
} else {
  gradient_calc <- readRDS(file = here("output", "calc_gr.rds"))
}

```

Classify each time point as either a surge/upswing or not based on established criteria.

```

gradient_classif <- gradient_calc %>%
  mutate(surge_pred = case_when(
    pred_gr >= surge_thresh & obs >= min_inc ~ TRUE,
    is.na(pred_gr) ~ NA,
    TRUE ~ FALSE
  )) %>% group_by(geo_value, model) %>%
  epi_slide(~{
    bef <- .x$surge_pred[1]
    focal <- .x$surge_pred[2]
    if (is.na(focal) | is.na(bef)){
      out <- NA
    } else {
      if (bef == FALSE & focal == TRUE){

```

```

      out <- TRUE
    } else if (bef == TRUE & focal == FALSE){
      out <- FALSE
    } else {
      out <- FALSE
    }
  }
  return(out)
}, n = 2 * 7 * 1, align = "right", new_col_name = "upswing_pred")

head(gradient_classif)

```

```

## # A tibble: 6 x 11
##   geo_value model      time_value population geo_type  obs surge upswing pred_gr
##   <chr>      <chr>      <date>          <dbl> <chr>    <dbl> <lgl> <lgl>    <dbl>
## 1 ak        CovidAna~ 2020-10-31      731545 state    2812 FALSE NA      NA
## 2 ak        CovidAna~ 2020-11-07      731545 state    2951 FALSE FALSE NA
## 3 ak        CovidAna~ 2020-11-14      731545 state    3846 FALSE FALSE NA
## 4 ak        CovidAna~ 2020-11-21      731545 state    4080 FALSE FALSE 0.346
## 5 ak        CovidAna~ 2020-11-28      731545 state    4225 FALSE FALSE 0.0266
## 6 ak        CovidAna~ 2020-12-05      731545 state    4829 FALSE FALSE -0.258
## # ... with 2 more variables: surge_pred <lgl>, upswing_pred <lgl>

```

Perform evaluation

```

surge_eval <- map_dfr(models, ~{
  misclass_surge <- gradient_classif %>% ungroup() %>%
    filter(model == .x) %>%
    filter(!is.na(surge_pred)) %>%
    summarise(misclass = mean(surge != surge_pred),
              sens = sum(surge & surge_pred)/sum(surge),
              spec = sum(!surge & !surge_pred)/sum(!surge))
  pmod <- pROC::roc(surge ~ pred_gr, data = gradient_classif %>% filter(model == .x),
                   subset = !is.na(surge_pred))
  misclass_surge <- mutate(misclass_surge, auc = round(as.numeric(pmod$auc),3), type = "surge")
  misclass_surge <- misclass_surge %>% mutate(model = .x)
})

upswing_eval <- map_dfr(models, ~{
  misclass_upswing <- gradient_classif %>% ungroup() %>%
    filter(model == .x) %>%
    filter(!is.na(upswing_pred)) %>%
    summarise(misclass = mean(upswing != upswing_pred),
              upswing_prev = mean(upswing),
              sens = sum(upswing & upswing_pred)/sum(upswing),
              spec = sum(!upswing & !upswing_pred)/sum(!upswing))
  pmod <- pROC::roc(upswing ~ pred_gr, data = gradient_classif %>% filter(model == .x),
                   subset = !is.na(upswing_pred))
  misclass_upswing <- mutate(misclass_upswing, auc = round(as.numeric(pmod$auc),3), type = "upswing")
  misclass_upswing <- misclass_upswing %>% mutate(model = .x)
})

```



```
saveRDS(upswing_eval, file = here("output", "upswing_eval.rds"))
saveRDS(surge_eval, file = here("output", "surge_eval.rds"))
knitr::kable(surge_eval)
```

misclass	sens	spec	auc	type	model
0.2283544	0.3076923	0.9700036	0.848	surge	COVIDhub-baseline
0.2375130	0.2826283	0.9575558	0.796	surge	RobertWalraven-ESG
0.2556164	0.2063179	0.9510808	0.542	surge	CovidAnalytics-DELPHI
0.1792767	0.5835498	0.9205539	0.848	surge	USC-SI_kJalpha
0.1951899	0.4167371	0.9707264	0.870	surge	COVIDhub-4_week_ensemble
0.1846753	0.5290043	0.9380334	0.842	surge	CU-nochange
0.1548223	0.7113885	0.8856873	0.880	surge	Karlen-pypm
0.1960526	0.4709507	0.9459459	0.836	surge	CU-scenario_low
0.1424390	0.5243446	0.9749340	0.871	surge	COVIDhub-ensemble

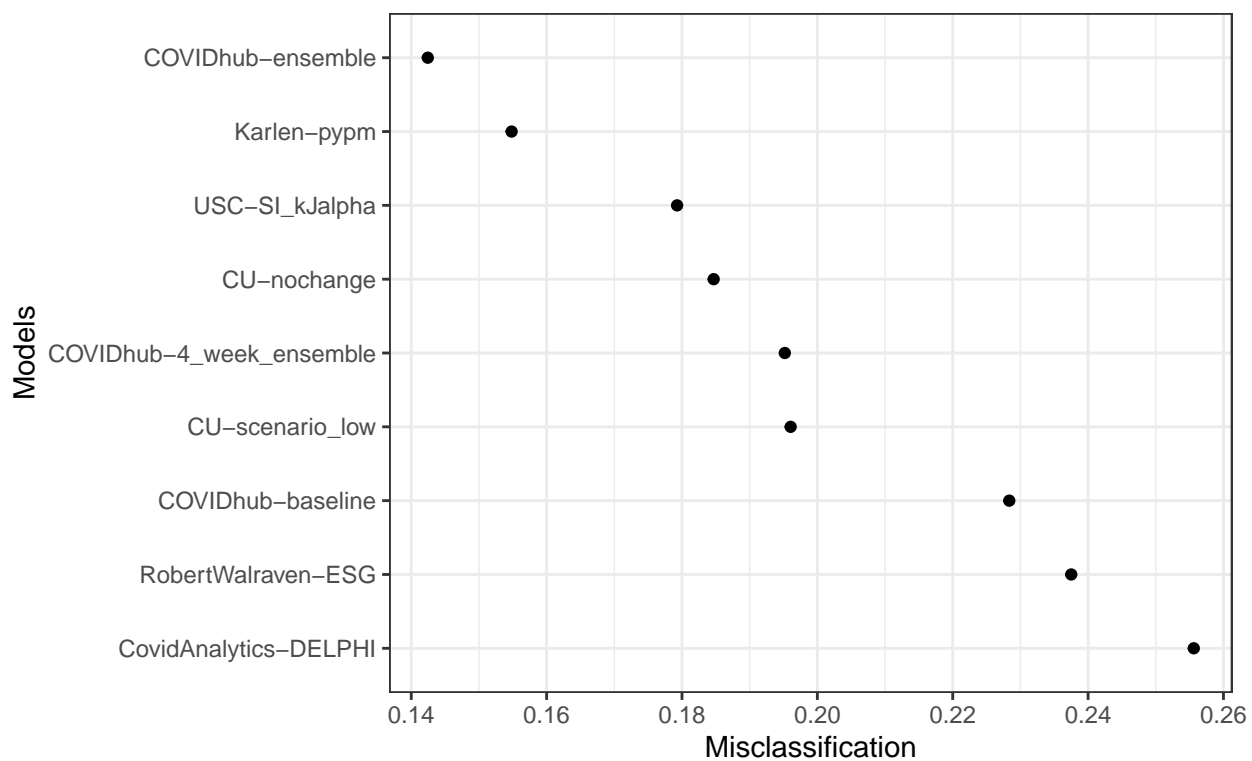
```
knitr::kable(upswing_eval)
```

misclass	upswing_prev	sens	spec	auc	type	model
0.0853846	0.0451282	0.0568182	0.9551557	0.518	upswing	COVIDhub-baseline
0.0826558	0.0436314	0.0062112	0.9589119	0.575	upswing	RobertWalraven-ESG
0.0751515	0.0463636	0.0065359	0.9694948	0.637	upswing	CovidAnalytics-DELPHI
0.1057692	0.0452183	0.1954023	0.9273272	0.559	upswing	USC-SI_kJalpha
0.0871795	0.0451282	0.0511364	0.9535446	0.501	upswing	COVIDhub-4_week_ensemble
0.1058667	0.0466667	0.1542857	0.9303497	0.541	upswing	CU-nochange
0.0846325	0.0378619	0.1176471	0.9467593	0.496	upswing	Karlen-pypm
0.1019178	0.0413699	0.0662252	0.9339811	0.502	upswing	CU-scenario_low
0.0660000	0.0320000	0.0156250	0.9643595	0.619	upswing	COVIDhub-ensemble

Visualize performance values

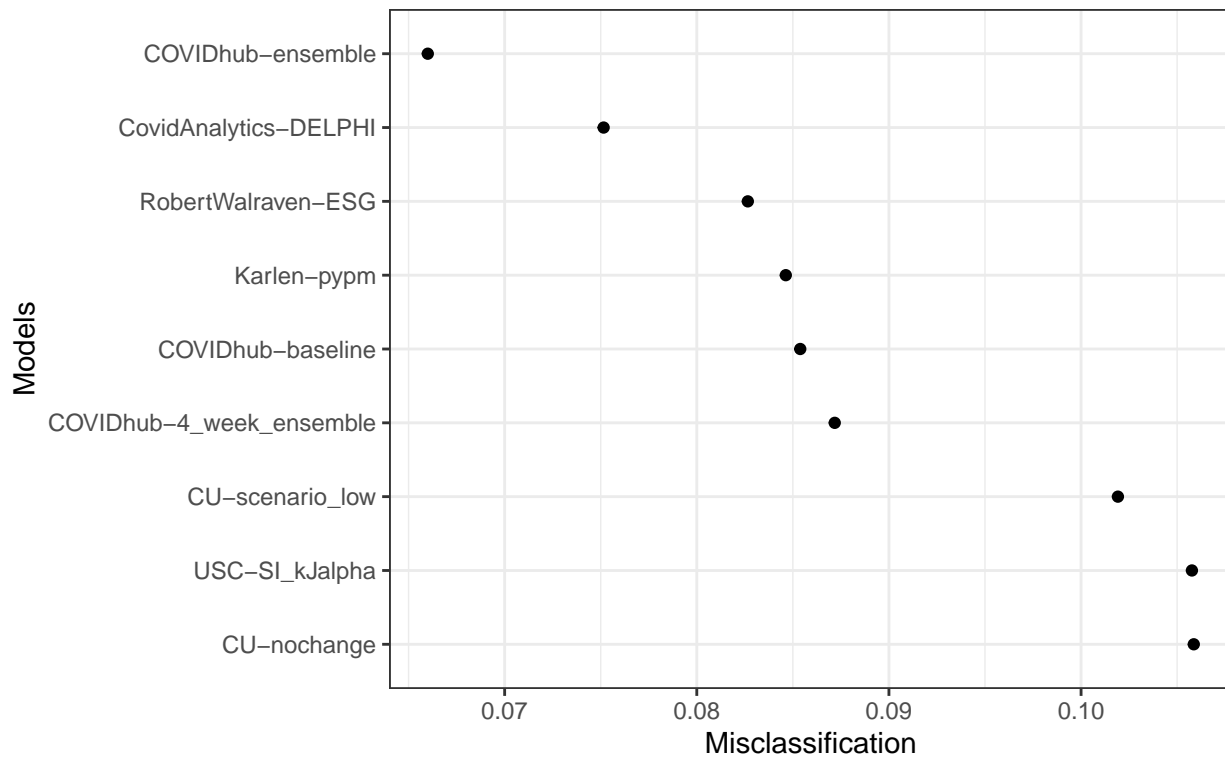
```
ggplot(surge_eval, aes(y = misclass, x = reorder(model, -misclass))) +
  coord_flip() + geom_point() +
  labs(x = "Models", y = "Misclassification", title = str_wrap("Classifying whether a week is 'surging'"))
```

## Classifying whether a week is 'surging' based on growth rate estimation



```
ggplot(upswing_eval, aes(y = misclass, x = reorder(model, -misclass))) +
  coord_flip() + geom_point() +
  labs(x = "Models", y = "Misclassification", title = str_wrap("Classifying whether a week is an upswing"))
```

## Classifying whether a week is an upswing or the beginning of the surge period



ROC curves for upswing and surge calculations

```
library(pals)
colors <- pals::glasbey(n = 9)

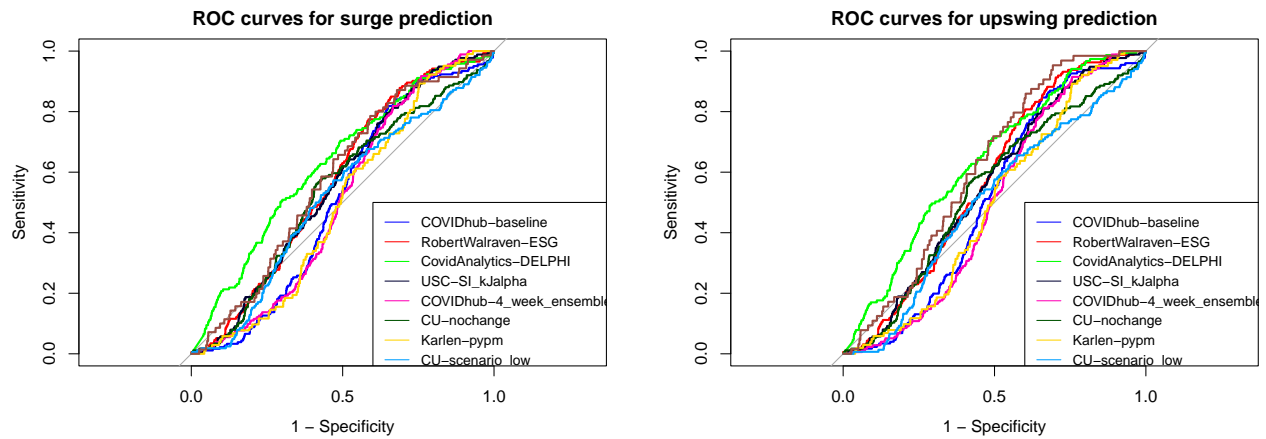
par(mfrow = c(1,2))

suppressMessages(mods <- map(models, ~{
  pROC::roc(upswing ~ pred_gr, data = gradient_classif %>% filter(model == .x),
    subset = !is.na(surge_pred), verbose = FALSE)
}))
names(mods) <- models
plot(mods[[1]], col = colors[1], legacy.axes = TRUE, main = "ROC curves for surge prediction")
for (i in 2:length(mods)){
  plot(mods[[i]], add = TRUE, col = colors[i])
}

legend(0.4,0.5,legend=models, col = colors, lty = 1, cex = 0.8)

suppressMessages(mods <- map(models, ~{
  pROC::roc(upswing ~ pred_gr, data = gradient_classif %>% filter(model == .x),
    subset = !is.na(upswing_pred), verbose = FALSE)
}))
names(mods) <- models
plot(mods[[1]], col = colors[1], legacy.axes = TRUE, main = "ROC curves for upswing prediction")
for (i in 2:length(mods)){
  plot(mods[[i]], add = TRUE, col = colors[i])
}
```

```
}
legend(0.4,0.5,legend=models, col = colors, lty = 1, cex = 0.8)
```

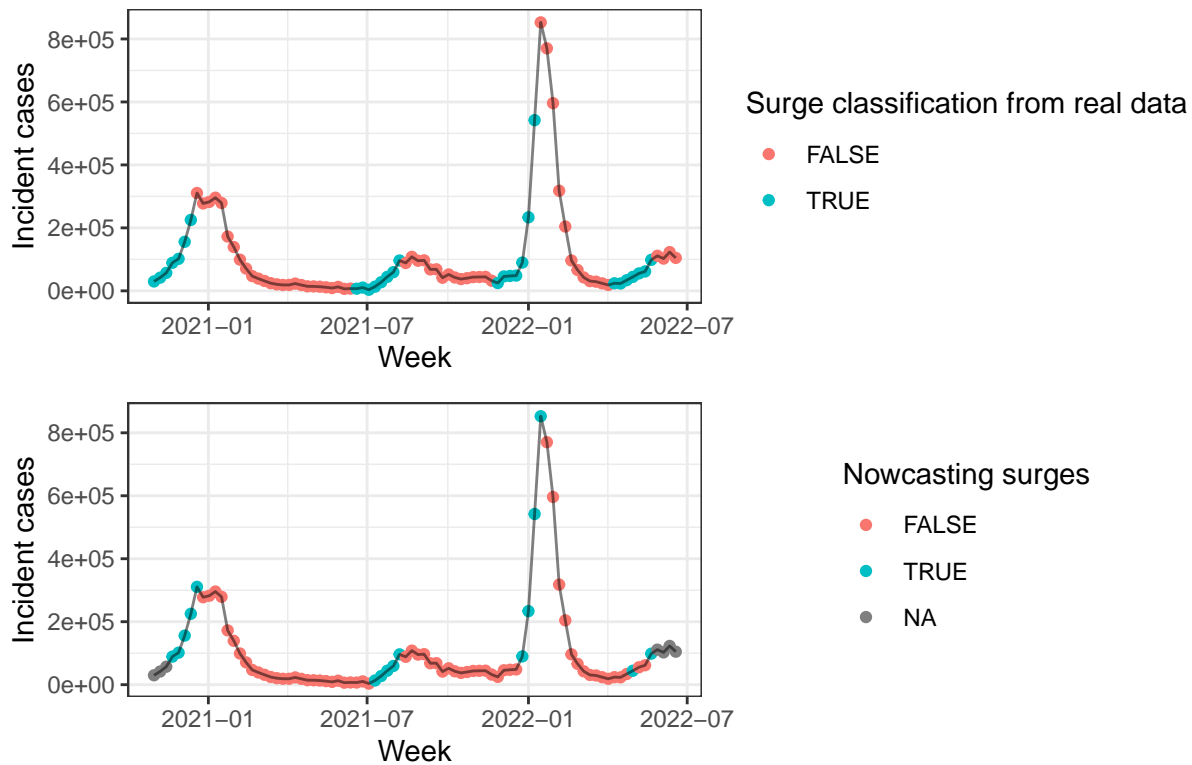


## Ensemble only evaluation

```
ensemble <- gradient_classif %>% filter(model == models[5])

p1 <- ggplot(ensemble %>% filter(geo_value == "ca"),
  aes(x = time_value, y = obs)) + geom_point(aes(col = surge)) + geom_line(alpha = 0.5) +
  labs(x = "Week", col = "Surge classification from real data", y = "Incident cases")
p2 <- ggplot(ensemble %>% filter(geo_value == "ca"), aes(x = time_value, y = obs)) +
  geom_point(aes(col = surge_pred)) +
  geom_line(alpha = 0.5) +
  labs(x = "Week", col = "Nowcasting surges", y = "Incident cases")
p1 / p2 + plot_annotation("Ensemble model nowcasting surges for the state of California")
```

## Ensemble model nowcasting surges for the state of California



Plotting incident cases and surge classification

```
p1 <- ggplot(ensemble %>% filter(geo_value == "ca"),
             aes(x = time_value, y = obs)) +
  geom_point(aes(col = upswing)) +
  geom_line(alpha = 0.5) +
  labs(x = "Week", col = "Upswing classification from real data", y = "Incident cases")
p2 <- ggplot(ensemble %>% filter(geo_value == "ca"), aes(x = time_value, y = obs)) +
  geom_point(aes(col = upswing_pred)) +
  geom_line(alpha = 0.5) +
  labs(x = "Week", col = "Nowcasting upswings", y = "Incident cases")

p1 / p2 + plot_annotation(title = "Ensemble model nowcasting upswings for the state of California")
```

Ensemble model nowcasting upswings for the state of California

