SemiMech

First we load the library.

The main function exported is called semimech:generate_samples() which takes a start argument, the beginning of the time series. dates a sequence of dates to forecast, region a flag for which geographic resolution to forecast (right now only US states), target a string ("hosp", "cases", "deaths") representing the target (right now set to hospitalizations) and inc_dat a data frame of disease incidence data (right now must be the result of a call to get hhs.)

Get the data!

```
library(stringr)
hhs <- get_hhs()</pre>
```

Define the forecast dates.

```
dates <- c(as.Date("2021-07-12"))
```

Get the samples.

```
samples <- semimech::generate_samples("2020-08-01",dates,"states",hhs)</pre>
```

Plot the samples

```
library(ggplot2)
median_mat <- matrix(NA,nrow=length(names(samples)),ncol=30)
upper_ci_mat <- matrix(NA,nrow=length(names(samples)),ncol=30)
lower_ci_mat <- matrix(NA,nrow=length(names(samples)),ncol=30)

row_idx <- 1
for (key in names(samples)){
    median_mat[row_idx,] <- colMeans(samples[[key]])
    upper_ci_mat[row_idx,] <- apply(samples[[key]],2,function(x){quantile(x,probs=.975)})
    lower_ci_mat[row_idx,] <- apply(samples[[key]],2,function(x){quantile(x,probs=.025)})

row_idx <- row_idx + 1
}

ci_mat_df <- data.frame(median_ = c(median_mat),upper_ci=c(upper_ci_mat),lower_ci=c(lower_ci_mat),st

p <- ggplot(ci_mat_df,aes(x=date,y=median_)) + geom_line() +
    geom_point(data=hhs[hhs$date > dates-30 & hhs$date < dates+30,],aes(x=as.Date(date),y=previous_day)

ggsave(filename=paste0("../figs/",dates[1],".png"),plot=p,height = 10,width = 12)</pre>
```

Generate submission file.

```
semimech::generate_submission_file(dates,samples,"UT-test","sub_files")
## Rows: 51
## Columns: 4
## $ fips
                               <int> 1, 2, 4, 5, 6, 8, 9, 10, 11, 12, 13, 15, 16, 17, 18, 19, 20~
## $ state_full <chr> "Alabama", "Alaska", "Arizona", "Arkansas", "California", "~
## $ state
                               <chr> "AL", "AK", "AZ", "AR", "CA", "CO", "CT", "DE", "DC", "FL",~
## $ alphacount <int> 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, ~
Let's do a full years worth of eval.
date seq <- seq(as.Date("2021-07-12"),as.Date("2021-07-12")+50*7,by="week")
for (date idx in 1:length(date seq)){
   samples <- semimech::generate_samples("2020-08-01",date_seq[date_idx],"states",hhs)</pre>
   semimech::generate_submission_file(date_seq[date_idx],samples,"UT-test","sub_files")
}
Let's evaluate with respect to baseline and ensemble. Note Before you evaluate you must copy
the files from sub_files and placed in the data_processed/UT-test folder in a cloned version of
https://github.com/reichlab/covid19-forecast-hub.
library(covidHubUtils)
fcasts <- load forecasts(</pre>
   source = "local hub repo",
   models =c("UT-test"),
   targets = c(paste0(rep(1:28), " day ahead inc hosp")),
   data_processed_subpath = "data-processed/",
   hub_repo_path = "/Users/gcg799/covid19-forecast-hub/",
   hub = "US")
fcasts_baseline <- load_forecasts(</pre>
   source = "local_hub_repo",
   models =c("COVIDhub-baseline","COVIDhub-ensemble"),date_window_size = 3,
   dates = unique(fcasts$forecast_date),locations = unique(fcasts$location),
   targets = c(pasteO(rep(1:28), " day ahead inc hosp")),
   data processed subpath = "data-processed/",
   hub_repo_path = "/Users/gcg799/covid19-forecast-hub/",
   hub = "US")
truth <- load truth(hub = "US")
truth <- truth[truth$target_variable == "inc hosp",]</pre>
scores <- score_forecasts(rbind(fcasts,fcasts_baseline),truth = truth,metrics = c("abs_error","wis",</pre>
scores_common_location <- unique(scores[scores*model == "UT-test",] $location)</pre>
scores_common_forecast_date <- unique(scores[scores$model == "UT-test",]$forecast_date)</pre>
\verb|summ_scores| <- \verb|scores| scores| 
                           scores$forecast_date %in% scores_common_forecast_date,] %>% group_by(model) %>% summar
print (summ_scores)
## # A tibble: 3 x 4
## model
```

wis abs_error

coverage_90

##		<chr></chr>	<dbl></dbl>	<dbl></dbl>	<dbl></dbl>
##	1	COVIDhub-baseline	0.906	54.0	78.8
##	2	COVIDhub-ensemble	0.875	34.5	52.9
##	3	UT-test	0.848	38.8	57.0