# 7 Appendix

**Note:** The source code can be found on Github in the repository found in the references. The primary files to consider are the source files found in the "final" directory, which will be refactored and tidied in an ongoing manner.

### 7.1 Data

- instant: record index
- dteday: date
- season: season (1:spring, 2:summer, 3:fall, 4:winter)
- yr: year (0:2011, 1:2012)
- mnth: month (1 to 12)
- hr: hour (0 to 23)
- holiday: weather day is holiday or not (extracted from http://dchr.dc.gov/page/holiday-schedule)
- weekday: day of the week
- workingday: if day is neither weekend nor holiday is 1, otherwise is 0.
- weather-sit:
  - 1: Clear, Few clouds, Partly cloudy, Partly cloudy
  - 2: Mist + Cloudy, Mist + Broken clouds, Mist + Few clouds, Mist
  - 3: Light Snow, Light Rain + Thunderstorm + Scattered clouds, Light Rain + Scattered clouds
  - 4: Heavy Rain + Ice Pallets + Thunderstorm + Mist, Snow + Fog
- temp: Normalized temperature in Celsius. The values are divided to 41 (max)
- atemp: Normalized feeling temperature in Celsius. The values are divided to 50 (max)
- hum: Normalized humidity. The values are divided to 100 (max)
- windspeed: Normalized wind speed. The values are divided to 67 (max)
- casual: count of casual users
- registered: count of registered users
- cnt: count of total rental bikes including both casual and registered

# 7.2 Wrangling

```
#======#
# Wrangling Function
#=======#
# Initial wrangling wrapper function
wrangle_init <- function(data, omit_NA = TRUE, omit_idx = TRUE){</pre>
  # Boolean variables (from int to logical type)
 data$holiday <- as.logical(data$holiday)</pre>
                                                   # 0 or 1
 data$workingday <- as.logical(data$workingday)</pre>
                                                 # 0 or 1
 # Other categorical variables (from int to factor type)
 data$season <- as.factor(data$season)</pre>
                                                  # 1 to 4
 data$yr <- as.factor(data$yr)</pre>
                                                  # 0 to 1
 data$mnth <- as.factor(data$mnth)</pre>
                                                  # 1 to 12
 data$weekday <- as.factor(data$weekday)</pre>
                                                 # 0 to 6
 data$weathersit <- as.factor(data$weathersit) # 1 to 4</pre>
 # Re-scale the normalized measurements
 data$temp <- data$temp * 41</pre>
 data$atemp <- data$atemp * 50</pre>
 data$hum <- data$hum * 100
 data$windspeed <- data$windspeed * 67</pre>
 # Change type of Dates (from char to Date type)
 data$dteday <- as.Date(data$dteday)</pre>
 # Remove NAs (if prompted) default value is TRUE
 if(omit_NA) { data <- na.omit(data) }</pre>
 # Remove instance column (if prompted) default value is TRUE
 if(omit_idx) { data <- data %>% select(-c("instant")) }
 # Observe christmas
 data$holiday[359] <- T; data$holiday[725] <- T</pre>
 # Return the wrangled dataset
 return(data)
}
#======#
# Subsetting
#----#
# Filter for the 2011 data
in_2011 <- function(data){ data[(data$dteday >= "2011-01-01" & data$dteday <= "2011-12-31"),] }
# Filter for the 2012 data
in_2012 \leftarrow function(data) \{ data[(data$dteday >= "2012-01-01" & data$dteday <= "2012-12-31"),] \}
#======#
   Wrangling
#======#
# Import and wrangle data
bike_day <- read.csv("bike-day.csv", header = T)</pre>
# Use the wrangling "wrapper" function to clean the data (fxn sourced from wrangle.R)
data <- wrangle_init(bike_day)</pre>
# Partition the 2011/2012 data
data2011 <- in_2011(data); data2012 <- in_2012(data)</pre>
```

## 7.3 Modeling

### 7.3.1 Model Building

```
#=======#
# Helper Function #
#======#
# A helper function that returns a formula in the "lm" syntax
# Takes predictors, a vector of variable name strings as an input
.parseFormula <- function(predictors, response = "cnt"){</pre>
 f <- as.formula(</pre>
   paste(response,
         paste(predictors, collapse = " + "),
         sep = " ~ "))
 return(f)
}
#========#
      Model "Builder" Functions
#=======#
# Given a vector of variable names, return a linear model fitting cnt
lm.tot <- function(varset, train_data = data2011){</pre>
 lm(formula = .parseFormula(predictors = varset, response = "cnt"), data = train_data)
# Given a vector of variable names, return a linear model fitting cas
lm.cas <- function(varset, train_data = data2011){</pre>
 lm(formula = .parseFormula(predictors = varset, response = "casual"), data = train data)
}
# Given a vector of variable names, return a linear model fitting reg
lm.reg <- function(varset, train_data = data2011){</pre>
 lm(formula = .parseFormula(predictors = varset, response = "registered"), data = train_data)
}
```

#### 7.3.2 Model Formulas

### 7.3.3 Growth-Adjusting Predictions

```
# Select some arbitrary model
model_example <- mod.cas.1.final
# Select a parameter estimate for g
g_hat <- 1.61
# Predictions without growth adjustment
preds <- predict(model_example, data2012)
# Predictions after growth adjustment
preds_adj <- preds * g_hat</pre>
```

## 7.4 Diagonostic Analysis

```
residual QQ <- function(model, title str = ""){</pre>
  # Plot parameters
  fsize0 <- 13; fsize1 <- 13; fsize2 <- 13
  # Plot
  df res <- data.frame(ep = model$residuals)</pre>
  df_res %>%
    ggplot(aes(sample = ep)) +
    geom_qq(alpha = 0.8, color = "palegreen3") +
    stat_qq_line(color = "seagreen") +
    labs(x = "Theoretical Quantile", y = "Sample Quantiles",
         title = paste("Normal Q-Q Plot", title_str, sep = ""))
}
residual_fitted <- function(model, title_str = ""){</pre>
  # Plot parameters
  fsize0 <- 13; fsize1 <- 13; fsize2 <- 13
  # Plot
  df_res <- data.frame(fitted_value = model$fitted.values, residual = model$residuals)</pre>
  df res %>%
    ggplot(aes(x = fitted_value, y = residual)) +
    geom_point(alpha = 0.7, color = "palegreen3") +
    stat_smooth(se = T, color = "forestgreen") +
    labs(x = "Fitted Value", y = "Residual",
         title = paste("Fitted vs Residual",title_str, sep = ""))
}
residual_histogram <- function(model, title_str = ""){</pre>
  # Plot parameters
  fsize0 <- 13; fsize1 <- 13; fsize2 <- 13
  epsilons <- unname(model$residuals)</pre>
  df_res <- data.frame(epsilon = epsilons)</pre>
  df_res %>%
    ggplot(aes(x = epsilon)) +
    geom_histogram(fill = "forestgreen", bins = 60) +
    labs(x = "Residual", title = paste("Residual Distribution", title_str, sep = ""))
}
residual_plots <- function(model, title_str){</pre>
  p1 <- model %>% residual_QQ(title_str)
  p2 <- model %>% residual_histogram(title_str)
 p3 <- model %>% residual_fitted(title_str)
  (p1 + p2 + p3)
```

### 7.5 Validation and Problemshooting

```
#
                            LOOCV
get_loocv_rmse = function(model) {
 loocv_rmse <- sqrt(mean((resid(model) / (1 - hatvalues(model))) ^ 2))</pre>
 # Round the numbers
 digits <- 2
 loocv_rmse <- round(loocv_rmse, digits)</pre>
 # Return the dataframe row
 return(data.frame('CV_rmse' = loocv_rmse))
}
get loocv rmse tot = function(cas, reg) {
 res_cas <- resid(cas) / (1 - hatvalues(cas))</pre>
 res_reg <- resid(reg) / (1 - hatvalues(reg))</pre>
 loocv_rmse <- sqrt(mean((res_cas + res_reg) ^ 2))</pre>
 # Round the numbers
 digits <- 2
 loocv_rmse <- round(loocv_rmse, digits)</pre>
 # Return the dataframe row
 return(data.frame('CV_rmse' = loocv_rmse))
}
Model Validation Functions
get_rmse <- function(y, y_hat, name='none'){</pre>
 rmse <- sqrt(mean((y - y hat)^2, na.rm = TRUE))</pre>
 normalized_rmse <- sqrt(mean(((y - y_hat)^2), na.rm = TRUE)) / sd(y)</pre>
 percentage_error <- mean(abs(y - y_hat) / y, na.rm = TRUE) * 100</pre>
 # Round the numbers
 digits <- 2
 rmse <- round(rmse, digits)</pre>
 normalized_rmse <- round(normalized_rmse, digits)</pre>
 percentage_error <- round(percentage_error, digits)</pre>
 # Return the dataframe row
 return(data.frame('name' = name, 'rmse' = rmse,
                  'nrmse' = normalized_rmse, 'prc_err' = percentage_error))
}
get_mod_eval_2011 <- function(cas, reg){</pre>
 ret1 <- rbind(get rmse(data2011$casual,
                                      predict(cas, data2011), '2011 cas'),
              get_rmse(data2011$registered, predict(reg, data2011), '2011 reg'),
              get_rmse(data2011$cnt,
                      predict(cas, data2011) + predict(reg, data2011), '2011 tot'))
 ret2 <- rbind(get loocv rmse(cas),</pre>
              get_loocv_rmse(reg),
              get_loocv_rmse_tot(cas, reg))
 return(cbind(ret1, ret2))
}
```

```
get_mod_eval <- function(cas, reg, data2011, data2012, scale_2012 = TRUE, include_2011 = F){</pre>
  if(scale_2012){G_FACTOR \leftarrow 0.608} else{G_FACTOR \leftarrow 1}
  if(include 2011){
    return(rbind(
      get_rmse(data2011$casual,
                                     predict(cas, data2011), '2011 cas'),
      get_rmse(data2011$registered, predict(reg, data2011), '2011 reg'),
      get_rmse(data2011$cnt,
                                     predict(cas, data2011) + predict(reg, data2011), '2011 tot'),
      get rmse(data2012$casual,
                                     predict(cas, data2012)/G FACTOR, '2012 cas'),
      get_rmse(data2012$registered, predict(reg, data2012)/G_FACTOR, '2012 reg'),
      get_rmse(data2012$cnt,
               predict(cas, data2012)/G_FACTOR + predict(reg, data2012)/G_FACTOR, '2012 tot')))
    }
  return(rbind(
    get_rmse(data2012$casual,
                                   predict(cas, data2012)/G_FACTOR, '2012 cas'),
    get_rmse(data2012$registered, predict(reg, data2012)/G_FACTOR, '2012 reg'),
    get_rmse(data2012$cnt,
             predict(cas, data2012)/G_FACTOR + predict(reg, data2012)/G_FACTOR, '2012 tot')))
get_mod_eval_tot_2011 <-function(tot){</pre>
 ret1 <- rbind(get_rmse(data2011$cnt, predict(tot, data2011), '2011 tot'))</pre>
  ret2 <- rbind(get loocv rmse(tot))</pre>
 return(cbind(ret1, ret2))
get_mod_eval_tot <-function(tot, data2011, data2012, scale_2012 = TRUE){</pre>
  if(scale_2012)\{G_FACTOR \leftarrow 0.608\} else\{G_FACTOR \leftarrow 1\}
    return(rbind(get_rmse(data2011$cnt, predict(tot, data2011), '2011 tot'),
                 get_rmse(data2012$cnt, predict(tot, data2012)/G_FACTOR, '2012 tot'),
                 get_loocv_rmse(tot)))
```

# 7.6 Prediction of the Yearly Growth Ratio

```
#----#
#
      Miscellaneous
#=======#
# Enumerate all the pairs in the lower-triangular matrix scheme
# In other words, (i < j or i - j < 0) so that day_i preceeds day_j
.unique_pairs_lower <- function(N){</pre>
  is <- do.call("c", purrr::map(1:N, function(i){rep(i,N)}))</pre>
  js \leftarrow rep(1:N, N)
  # Helper function: selects elements only if they are upper triangular
  .LowerTri \leftarrow function(i, j){if(i > j) { c(i = i, j = j) }}
 pairs <- do.call("rbind", purrr::map2(is, js, .f = .LowerTri))</pre>
 data.frame(pairs)
}
#=======#
     Dataframe Wrapper Functions
#======#
# Given a dataframe of loss values between all possible day pairs
# and a set of idx and loss bounds, compute the q estimates
get_df_param <- function(df_loss, idx_bds, loss_bds){</pre>
  # Compute the index pairs
  MIP <- .unique_pairs_lower(length(idx_bds))</pre>
  # Use helper function to compute the g estimates for each set of bounds
  .helper <- function(k, df_loss = df_loss){</pre>
   idx <- MIP$i[k]; loss <- MIP$j[k]</pre>
   results <- g_estimate(df_loss, idx, loss)
   data.frame(idx_bd = idx_bds[idx], loss_bd = loss_bds[loss], g = results[[1]], n = results[[2]])
  # Run helper over 1,...,n where n is the number of pairs surviving bound cutoff
  df_param <- map_dfr(1:nrow(MIP), .helper, df_loss)</pre>
  # Return the estimates and number of pairs used for each bound pair
 return(df_param)
}
get_df_loss <- function(data2011){</pre>
  # Obtain the continous variables and holiday
  data <- data2011 %>% select(dteday, holiday, atemp, hum, windspeed, "cnt")
  # Normalize the continous variables
  data <- data %>% mutate_at(c("atemp", "hum", "windspeed"), ~(scale(.) %>% as.vector))
  # Take out holidays (to avoid needing to account for its effect, omits only few days of data)
  data <- data %>% filter(!holiday)
  # Enumerate all the possible index pairs (for use in purrr::map2_dfr)
  pairs <- .unique_pairs_lower(nrow(data))</pre>
  # Helper function that takes two indices and returns the loss between the days at those indices
  .helper <- function(i,j){data.frame(i = i, j = j, loss_ij = loss_ij(i,j))}</pre>
  # Compute the loss function exhaustively for every possible (lower-triangle) pair
  df_loss <- map2_dfr(pairs[["i"]], pairs[["j"]], .f = .helper)</pre>
  # Add the index difference column (useful for determining space between days)
  df_loss <- df_loss %>% mutate(idx_diff = i - j)
  # Return the exhaustive dataset of loss values for every unique day ordered pair
  return(df_loss)
}
```

```
#----#
      Loss Function
#======#
# Given two rows (days) from the data, compute the loss function
loss <- function(day0, day1){</pre>
 atemp_diff <- day0[["atemp"]] - day1[["atemp"]]</pre>
 wind_diff <- day0[["windspeed"]] - day1[["windspeed"]]</pre>
 hum_diff <- day0[["hum"]] - day1[["hum"]]</pre>
 norm(as.matrix(c(4*atemp_diff, wind_diff, hum_diff)))
}
# Given two row indices, compute the loss function between those two days
loss_ij <- function(i, j){</pre>
 day0 <- data[i,]; day1 <- data[j,]</pre>
 loss(day0, day1)
}
growth_ratio <- function(i, j){ data2011$cnt[i]/data2011$cnt[j] }</pre>
g_estimate <- function(df_loss, idx_bound, loss_bound){</pre>
 # Compute the g-estimate after filtering through bounds
 df_loss <- df_loss %>%
   filter(idx_diff > idx_bound) %>%
   filter(loss_ij < loss_bound) %>%
   mutate(g = growth_ratio(i, j))
 # Return the mean (q^{\hat{}}) and number of pairs used (sample size n)
 list(mean(df_loss$g), nrow(df_loss))
}
#======#
     Loss Fxn Technique Plots
#======#
g_plot <- function(df_param){</pre>
 # Filter bounds to obtain a good window frame
 df_param <- df_param %>% filter(idx_bd > 10, loss_bd < 5)</pre>
 # Create the proportion of maximum pairs variable (for alpha)
 df_param <- df_param %>% mutate(perc_pairs_omitted = 1 - n/max(df_param$n))
 # Scatterplot
 df_param %>%
   ggplot() +
   geom_point(aes(color = idx_bd, x = loss_bd, y = g, alpha = perc_pairs_omitted)) +
   geom_abline(slope = 0, intercept = 3.35) +
   theme(legend.position = "bottom") +
   labs(x = "Loss Upper Bound", y = "g", title = "Growth Factor Esimates by Loss Upper Bound") +
   #scale_alpha(quide = "none") +
   scale_color_gradient(low = "palevioletred2", high = "seagreen3")
}
```

```
#----#
     Window Technique
#----#
window_g <- function(w, no_outlier = T){</pre>
 # Compute the indices of the first and last w days
 lower_idx <- 1:w</pre>
 upper_idx <- 366 - (w:1)
 # Obtain the first and last w days
 last_w <- data2011[upper_idx,]</pre>
 first_w <- data2011[lower_idx,]</pre>
 # Since Jan-3 is environmentally different (by loss function value), we remove it
 if(no_outlier){ first_w <- first_w %% filter(dteday != "2011-1-3") }</pre>
 # Compute and return the difference in means between the first and last w days
 return(mean(last_w$cnt)/mean(first_w$cnt))
}
#======#
     Window Technique Plots
#----#
plot_window <- function(tbl_window){</pre>
 # Plot parameters
 col0 <- "steelblue3"</pre>
 # Scatterplot
 tbl_window %>%
   ggplot(aes(w, g_w)) +
   geom_point(color = col0) +
   geom_line(color = col0) +
   geom_vline(xintercept = 6, color = "red") +
   labs(title = "Growth Factor Estimates by Window Size")
}
```

```
#-----#
#-----#
                   Primary Code Script
#-----#
#-----#
# Control flow for avoiding computational/time waste
recompute <- F
if(recompute){
 # Obtain the exhaustive dataset of loss values for every unique day ordered pair
 df_loss <- data_2011 %>% get_df_loss
 # Write CSV to avoid future recomputation
 write.csv(df_loss, "df_loss.csv", row.names = F)
} else{
 df_loss <- read.csv("df_loss.csv")</pre>
#========#
# Estimate Performance over Bound Paramater Space
# Set a sequence of index bounds
idx_bds \leftarrow seq(122, 365, 8)
# Set a sequence of loss bounds: more critical
loss_bds \leftarrow seq(1, 4, 0.10)
# Obtain the parameter dataframe of g estimates
df_param <- df_loss %>% get_df_param(idx_bds, loss_bds)
# Plot the estimates over bound parameter space
plot_g <- df_param %>% g_plot()
#======#
# Window Technique #
#======#
\# Obtain the table of g esimates by window parameter w
tbl_window <- purrr::map_dfr(1:20, function(w){data.frame(w = w, g_w = window_g(w))})
# Plot the values
plot_w <- plot_window(tbl_window)</pre>
```