

# Code Appendix

Taqi

12/10/2021

## Appendix

### Wrangling

```
#=====#
#      Wrangling Function      #
#=====#
# Initial wrangling wrapper function
wrangle_init <- function(data, omit_NA = TRUE, omit_idx = TRUE){
  # Boolean variables (from int to logical type)
  data$holiday <- as.logical(data$holiday)      # 0 or 1
  data$workingday <- as.logical(data$workingday) # 0 or 1
  # Other categorical variables (from int to factor type)
  data$season <- as.factor(data$season)         # 1 to 4
  data$yr <- as.factor(data$yr)                 # 0 to 1
  data$mnth <- as.factor(data$mnth)             # 1 to 12
  data$weekday <- as.factor(data$weekday)       # 0 to 6
  data$weathersit <- as.factor(data$weathersit)   # 1 to 4
  # Re-scale the normalized measurements
  data$temp <- data$temp * 41
  data$atemp <- data$atemp * 50
  data$hum <- data$hum * 100
  data$windspeed <- data$windspeed * 67
  # Change type of Dates (from char to Date type)
  data$dteday <- as.Date(data$dteday)
  # Remove NAs (if prompted) default value is TRUE
  if(omit_NA) { data <- na.omit(data) }
  # 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
  # Return the wrangled dataset
  return(data)
}
#=====#
#      Subsetting      #
#=====#
# Filter for the 2011 data
in_2011 <- function(data){
  return(data[(data$dteday >= "2011-01-01" & data$dteday <= "2011-12-31"),])
}
# Filter for the 2012 data
```

```

in_2012 <- function(data){
  return(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)
# Use the wrangling "wrapper" function to clean the data (fxn sourced from wrangle.R)
data <- wrangle_init(bike_day)
# Partition the 2011/2012 data
data2011 <- in_2011(data); data2012 <- in_2012(data)

```

## Modeling

### 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"){
  f <- as.formula(
    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){
  model <- lm(formula = .parseFormula(predictors = varset, response = "cnt"), data = train_data)
  model
}

# Given a vector of variable names, return a linear model fitting cas
lm.cas <- function(varset, train_data = data2011){
  model <- lm(formula = .parseFormula(predictors = varset, response = "casual"), data = train_data)
  model
}

# Given a vector of variable names, return a linear model fitting reg
lm.reg <- function(varset, train_data = data2011){
  model <- lm(formula = .parseFormula(predictors = varset, response = "registered"), data = train_data)
  model
}

```

### Model Formulas

```

# Initial models
mod.cas.1.final <- lm.cas(c("workingday", "weathersit", "atemp"))

```

```

mod.reg.1.final <- lm.reg(c("workingday", "weathersit", "atemp", "I(atemp^2)"))
mod.tot.1.final <- lm.tot(c("workingday", "weathersit", "atemp", "I(atemp^2)"))
# Final models
mod.cas.2.final <- lm.cas(c("holiday", "season:weathersit", "season:workingday:atemp"))
mod.reg.2.final <- lm.reg(c("holiday", "season:weathersit",
                           "season:workingday:atemp", "season:workingday:I(atemp^2)"))

```

## 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

```

## Diagnostic Analysis

```

residual_QQ <- function(model, title_str = ""){
  # Plot parameters
  fsize0 <- 13; fsize1 <- 13; fsize2 <- 13
  # Plot
  df_res <- data.frame(ep = model$residuals)
  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))
}

residual_fitted <- function(model, title_str = ""){
  # Plot parameters
  fsize0 <- 13; fsize1 <- 13; fsize2 <- 13
  # Plot
  df_res <- data.frame(fitted_value = model$fitted.values, residual = model$residuals)
  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 = ""){
  # Plot parameters
  fsize0 <- 13; fsize1 <- 13; fsize2 <- 13
  # Plot
  epsilons <- unname(model$residuals)
  df_res <- data.frame(epsilon = epsilons)
  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){
  p1 <- model %>% residual_QQ(title_str)
  p2 <- model %>% residual_histogram(title_str)
  p3 <- model %>% residual_fitted(title_str)
  (p1 + p2 + p3)
}

```

## Validation and Problemshooting

```

#####
#                               LOOCV
#####

```

```

get_loocv_rmse = function(model) {
  loocv_rmse <- sqrt(mean((resid(model) / (1 - hatvalues(model))) ^ 2))
  # Round the numbers
  digits <- 2
  loocv_rmse <- round(loocv_rmse, digits)
  # 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))
  res_reg <- resid(reg) / (1 - hatvalues(reg))
  loocv_rmse <- sqrt(mean((res_cas + res_reg) ^ 2))
  # Round the numbers
  digits <- 2
  loocv_rmse <- round(loocv_rmse, digits)
  # Return the dataframe row
  return(data.frame('CV_rmse' = loocv_rmse))
}

```

```

#####
#                               Model Validation Functions
#####

```

```

get_rmse <- function(y, y_hat, name='none'){
  rmse <- sqrt(mean((y - y_hat)^2, na.rm = TRUE))
  normalized_rmse <- sqrt(mean(((y - y_hat)^2), na.rm = TRUE)) / sd(y)
  percentage_error <- mean(abs(y - y_hat) / y, na.rm = TRUE) * 100
  # Round the numbers
  digits <- 2
  rmse <- round(rmse, digits)
  normalized_rmse <- round(normalized_rmse, digits)
  percentage_error <- round(percentage_error, digits)
  # Return the dataframe row
  return(data.frame('name' = name, 'rmse' = rmse, 'nrmse' = normalized_rmse, 'prc_err' = percentage_err
}

```

```

get_mod_eval_2011 <- function(cas, reg){
  ret1 <- rbind(get_rmse(data2011$casual,      predict(cas, data2011), '2011 casual'),

```

```

        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),
              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){
  if(scale_2012){G_FACTOR <- 0.608} else{G_FACTOR <- 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,
    )
    )
  }
  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){
  ret1 <- rbind(get_rmse(data2011$cnt, predict(tot, data2011), '2011 tot'))
  ret2 <- rbind(get_loocv_rmse(tot))
  return(cbind(ret1, ret2))
}

get_mod_eval_tot <-function(tot, data2011, data2012, scale_2012 = TRUE){
  if(scale_2012){G_FACTOR <- 0.608} else{G_FACTOR <- 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)))
}

```

## 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){
  is <- do.call("c", purrr::map(1:N, function(i){rep(i,N)}))
  js <- rep(1:N, N)
  # Helper function: selects elements only if they are upper triangular
  .LowerTri <- function(i, j){if(i > j) { c(i = i, j = j) }}
  pairs <- do.call("rbind", purrr::map2(is, js, .f = .LowerTri))
  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 g estimates
get_df_param <- function(df_loss, idx_bds, loss_bds){
  # Compute the index pairs
  MIP <- .unique_pairs_lower(length(idx_bds))
  # Use helper function to compute the g estimates for each set of bounds
  .helper <- function(k, df_loss = df_loss){
    idx <- MIP$i[k]; loss <- MIP$j[k]
    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)
  # Return the estimates and number of pairs used for each bound pair
  return(df_param)
}

get_df_loss <- function(data2011){
  # 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))
  # 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))}
  # Compute the loss function exhaustively for every possible (lower-triangle) pair
  df_loss <- map2_dfr(pairs[["i"]], pairs[["j"]], .f = .helper)
  # 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){
  atemp_diff <- day0[["atemp"]] - day1[["atemp"]]
  wind_diff <- day0[["windspeed"]] - day1[["windspeed"]]
  hum_diff <- day0[["hum"]] - day1[["hum"]]
  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){
  day0 <- data[i,]; day1 <- data[j,]
  loss(day0, day1)
}

growth_ratio <- function(i, j){ data2011$cnt[i]/data2011$cnt[j] }

g_estimate <- function(df_loss, idx_bound, loss_bound){
  # 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 (g) 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){
  # Filter bounds to obtain a good window frame
  df_param <- df_param %>% filter(idx_bd > 10, loss_bd < 5)
  # 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 Estimates by Loss Upper Bound") +
    #scale_alpha(guide = "none") +
    scale_color_gradient(low = "palevioletred2", high = "seagreen3")
}

#####
#####

#####
#      Window Technique      #
#####

window_g <- function(w, no_outlier = T){
  # Compute the indices of the first and last w days
  lower_idx <- 1:w
  upper_idx <- 366 - (w:1)
  # Obtain the first and last w days

```

```

last_w <- data2011[upper_idx,]
first_w <- data2011[lower_idx,]
# 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") }
# 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){
  # Plot parameters
  col0 <- "steelblue3"
  # 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")
}

#####
#      Estimate Performance over Bound Paramater Space
#####

# Set a sequence of index bounds
idx_bds <- seq(122, 365, 8)
# Set a sequence of loss bounds: more critical
loss_bds <- 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 estimates 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)

#####
#-----#
#####

```