## AlexBoffa-MovieLens-Project-Harvard.R

## YogaI5

## 2020-03-18

```
###################################
# Create edx set, validation set
###################################
# Note: this process took my computer about 10 min create edx and validation sets.
if(!require(tidyverse)) install.packages("tidyverse", repos="http://cran.us.r-project.org")
## Loading required package: tidyverse
## -- Attaching packages ------ tidyverse 1.3.0 -
## v ggplot2 3.3.0
                   v purrr
                              0.3.3
## v tibble 2.1.3
                   v dplyr 0.8.5
## v tidyr 1.0.2
                    v stringr 1.4.0
## v readr
          1.3.1
                    v forcats 0.5.0
## Warning: package 'ggplot2' was built under R version 3.6.3
## Warning: package 'dplyr' was built under R version 3.6.3
## Warning: package 'forcats' was built under R version 3.6.3
## -- Conflicts ----- tidyverse_conflicts() -
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                   masks stats::lag()
if(!require(caret)) install.packages("caret", repos="http://cran.us.r-project.org")
## Loading required package: caret
## Warning: package 'caret' was built under R version 3.6.3
## Loading required package: lattice
## Attaching package: 'caret'
## The following object is masked from 'package:purrr':
##
      lift
if(!require(data.table)) install.packages("data.table", repos="http://cran.us.r-project.org")
## Loading required package: data.table
## Warning: package 'data.table' was built under R version 3.6.3
## Attaching package: 'data.table'
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## The following objects are masked from 'package:dplyr':
##
##
       between, first, last
## The following object is masked from 'package:purrr':
##
##
       transpose
# MovieLens 10M dataset:
# https://grouplens.org/datasets/movielens/10m/
# http://files.grouplens.org/datasets/movielens/ml-10m.zip
dl <- tempfile()</pre>
download.file("http://files.grouplens.org/datasets/movielens/ml-10m.zip", dl)
ratings <- fread(text = gsub("::", "\t", readLines(unzip(dl, "ml-10M100K/ratings.dat"))),</pre>
                      col.names = c("userId", "movieId", "rating", "timestamp"))
movies <- str_split_fixed(readLines(unzip(dl, "ml-10M100K/movies.dat")), "\\::", 3)</pre>
colnames(movies) <- c("movieId", "title", "genres")</pre>
movies <- as.data.frame(movies) %>% mutate(movieId = as.numeric(levels(movieId))[movieId],
                                           title = as.character(title),
                                           genres = as.character(genres))
movielens <- left_join(ratings, movies, by = "movieId")</pre>
# Validation set will be 10% of MovieLens data
set.seed(1, sample.kind="Rounding")
## Warning in set.seed(1, sample.kind = "Rounding"): non-uniform 'Rounding' sampler
# if using R 3.5 or earlier, use `set.seed(1)` instead
test_index <- createDataPartition(y = movielens$rating, times = 1, p = 0.1, list = FALSE)
edx <- movielens[-test_index,]</pre>
temp <- movielens[test_index,]</pre>
# Make sure userId and movieId in validation set are also in edx set
validation <- temp %>%
  semi_join(edx, by = "movieId") %>%
  semi join(edx, by = "userId")
# Add rows removed from validation set back into edx set
removed <- anti_join(temp, validation)</pre>
## Joining, by = c("userId", "movieId", "rating", "timestamp", "title", "genres")
edx <- rbind(edx, removed)</pre>
rm(dl, ratings, movies, test_index, temp, movielens, removed)
##### Starting MovieLens Recomendation System Project
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# Install all needed libraries if they are not present
if(!require(tidyverse)) install.packages("tidyverse")
if(!require(tidyr)) install.packages("tidyr")
if(!require(stringr)) install.packages("stringr")
if(!require(forcats)) install.packages("forcats")
if(!require(ggplot2)) install.packages("ggplot2")
if(!require(kableExtra)) install.packages("kableExtra")
## Loading required package: kableExtra
## Warning: package 'kableExtra' was built under R version 3.6.3
## Attaching package: 'kableExtra'
## The following object is masked from 'package:dplyr':
##
      group_rows
# Loading all needed libraries
library(dslabs)
library(caret)
library(dplyr)
library(tidyverse)
library(kableExtra)
library(tidyr)
library(stringr)
library(forcats)
library(ggplot2)
# This is the RMSE function that will give us the scores found with our models.
# Our goal is obtain a RMSE < 0.86490
RMSE <- function(true_ratings, predicted_ratings) {</pre>
  sqrt(mean((true_ratings - predicted_ratings)^2))
}
# We start computing our models here #
##############################
### Starting Naive model ###
# Calculating "just the average" of all movies
mu <- mean(edx$rating)</pre>
\# Calculating the RMSE on the validation set
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naive_rmse <- RMSE(validation$rating, mu)</pre>
# Creating a results dataframe that will contains all RMSE results.
# Here we insert our first RMSE.
rmse_results <- data.frame(method = "Just the average", RMSE = naive_rmse)</pre>
### Starting Movie Effect Model ###
# Calculating the average by movie
movie_avgs <- edx %>%
   group_by(movieId) %>%
   summarize(b_i = mean(rating - mu))
# Computing the predicted ratings on validation dataset
predicted_ratings <- mu + validation %>%
  left_join(movie_avgs, by='movieId') %>%
   .$b i
# Computing Movie effect model
model_1_rmse <- RMSE(predicted_ratings, validation$rating)</pre>
# Adding the results to the rmse_results table
rmse_results <- bind_rows(rmse_results,</pre>
                         data.frame(method = "Movie Effect Model",
                                   RMSE = model_1_rmse ))
## Warning in bind_rows_(x, .id): Unequal factor levels: coercing to character
## Warning in bind_rows_(x, .id): binding character and factor vector, coercing
## into character vector
## Warning in bind_rows_(x, .id): binding character and factor vector, coercing
## into character vector
### Starting Movie + User Effect Model ###
# Calculating the average by user
user_avgs <- edx %>%
  left_join(movie_avgs, by='movieId') %>%
  group_by(userId) %>%
   summarize(b_u = mean(rating - mu - b_i))
\# Computing the predicted ratings on validation dataset
predicted_ratings <- validation %>%
  left_join(movie_avgs, by='movieId') %>%
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left_join(user_avgs, by='userId') %>%
  mutate(pred = mu + b_i + b_u) %>%
   .$pred
model_2_rmse <- RMSE(predicted_ratings, validation$rating)</pre>
# Adding the results to the results dataset
rmse_results <- bind_rows(rmse_results,</pre>
                        data.frame(method = "Movie + User Effects Model",
                                 RMSE = model_2_rmse ))
## Warning in bind_rows_(x, .id): binding character and factor vector, coercing
## into character vector
# Starting Regularization of our models.
# Regularizing Movie + User Effect Model
# Computing the predicted ratings on validation dataset using different values of lambda
\# b_i is the Movie effect and b_u is User effect.
# Lambda is a tuning parameter.
# We are using cross-validation to choose the best lambda that minimize our RMSE.
lambdas \leftarrow seq(0, 10, 0.25)
# function rmses calculate predictions with several lambdas
rmses <- sapply(lambdas, function(l) {</pre>
  # Calculating the average by movie
  b_i <- edx %>%
     group_by(movieId) %>%
     summarize(b_i = sum(rating - mu) / (n() + 1))
   # Calculating the average by user
  b_u <- edx %>%
     left_join(b_i, by="movieId") %>%
     group_by(userId) %>%
     summarize(b_u = sum(rating - b_i - mu) / (n() + 1))
   # Computing the predicted ratings on validation dataset
  predicted_ratings <- validation %>%
     left join(b i, by = 'movieId') %>%
     left_join(b_u, by = "userId") %>%
     mutate(pred = mu + b_i + b_u) %>%
     .$pred
   # Predicting the RMSE on the validation set
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return(RMSE(predicted_ratings, validation$rating))
})
# Getting the best lambda value that minimize the RMSE on reg movie + user effects model
lambda <- lambdas[which.min(rmses)]</pre>
# We know that our best RMSE is given by: RMSE = min(rmses),
# but as a purpose of clarification,
# we compute again our estimate with best lambda found:
# Compute regularized estimates of b_i using best lambda
movie_avgs_reg <- edx %>%
   group_by(movieId) %>%
   summarize(b_i = sum(rating - mu) / (n() + lambda), n_i = n())
# Compute regularized estimates of b_u using best lambda
user_avgs_reg <- edx %>%
   left_join(movie_avgs_reg, by ='movieId') %>%
   group_by(userId) %>%
   summarize(b_u = sum(rating - mu - b_i) / (n() + lambda), n_u = n())
# Predict ratings
predicted_ratings <- validation %>%
   left_join(movie_avgs_reg, by = 'movieId') %>%
   left_join(user_avgs_reg, by = 'userId') %>%
   mutate(pred = mu + b_i + b_u) %>%
   .$pred
# Predicting the RMSE on the validation set
model_3_rmse <- RMSE(predicted_ratings, validation$rating)</pre>
# Adding the results to the rmse_results dataset
rmse_results <- bind_rows(rmse_results,</pre>
                       data.frame(method = "Regularized Movie + User Effects Model",
                                  RMSE = model_3_rmse ))
## Warning in bind_rows_(x, .id): binding character and factor vector, coercing
## into character vector
rmse_results
##
                                      method
                                                  RMSE
## 1
                           Just the average 1.0612018
## 2
                         Movie Effect Model 0.9439087
                 Movie + User Effects Model 0.8653488
## 4 Regularized Movie + User Effects Model 0.8648170
```

```
\# Remmember, our goal is obtain a RMSE < 0.86490
# And voila, running this project in R on my computer,
# and just as info, I have obtained these rmse_results,
# where the "Regularized Movie + User Effects Model"
# exceeded the goal:
#
                        method
                                  RMSE
#
#1
                 Just the average 1.0612018
#2
               Movie Effect Model 0.9439087
#3 Movie + User Effects Model 0.8653488
#4 Regularized Movie + User Effects Model 0.8648170
# rmse_results %>% knitr::kable()
```