

ANA 515: Assignment 4

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1. Discuss the business problem/goal

The goals of this project (DataFlair (2023)) is to create a recommendation engine that suggests films that appeal to users based on their preferences and browsing history. The recommendations are powered by a Machine Learning algorithm; specifically an Item Based Collaborative Filter.

2. Identify where the dataset was retrieved from

The MovieLens dataset was retrieved from

<https://drive.google.com/file/d/1Dn1BZD3YxgBQJSIjbfNnmCF1DW2jdQGD/view>
which has two csv files – movies.csv and ratings.csv (DataFlair (2023)).

3. Identify the code that imported and saved your dataset in R

```
# Reading in the dataset, as downloaded from the links provided at  
# https://data-flair.training/blogs/data-science-r-movie-recommendation/  
# and saved into personal GitHub space:  
movie_data <- read_csv(  
  "https://raw.githubusercontent.com/dr3am05/Source/main/movies.csv"  
)  
  
## Rows: 10329 Columns: 3  
## -- Column specification -----  
## Delimiter: ","  
## chr (2): title, genres  
## dbl (1): movieId  
##  
## i Use `spec()` to retrieve the full column specification for this data.  
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.  
rating_data <- read_csv(  
  "https://raw.githubusercontent.com/dr3am05/Source/main/ratings.csv"  
)  
  
## Rows: 105339 Columns: 4  
## -- Column specification -----  
## Delimiter: ","  
## dbl (4): userId, movieId, rating, timestamp  
##  
## i Use `spec()` to retrieve the full column specification for this data.  
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.
```

4. Describe your data set

This dataset consists of 105339 ratings applied over 10329 movies. Movie data has 10329 rows with 3 columns, named `movieId`, `title` and `genres`. `movieId` is an integer type while `title` and `genres` are character type. Rating data has 105339 rows with 4 columns, named `userId`, `movieId`, `rating` and `timestamp`.

Below is a summary of our movie data and ratings data by each variable.

```
summary(movie_data)
```

```
##      movieId      title      genres
## Min.   :    1  Length:10329  Length:10329
## 1st Qu.: 3240  Class :character  Class :character
## Median : 7088  Mode  :character  Mode  :character
## Mean   : 31924
## 3rd Qu.: 59900
## Max.   :149532
```

```
summary(rating_data)
```

```
##      userId      movieId      rating      timestamp
## Min.   :  1.0  Min.   :    1  Min.   :0.500  Min.   :8.286e+08
## 1st Qu.:192.0  1st Qu.: 1073  1st Qu.:3.000  1st Qu.:9.711e+08
## Median :383.0  Median : 2497  Median :3.500  Median :1.115e+09
## Mean   :364.9  Mean   :13381  Mean   :3.517  Mean   :1.130e+09
## 3rd Qu.:557.0  3rd Qu.: 5991  3rd Qu.:4.000  3rd Qu.:1.275e+09
## Max.   :668.0  Max.   :149532  Max.   :5.000  Max.   :1.452e+09
```

The movie data has 0 missing values; and the ratings data has 0 missing values.

5. Discuss any data preparation, missing values and errors

As discussed in 4, there are no missing values in both datasets

Because Machine Learning needs data represented in matrix forms, we need to prepare our datasets into that format.

First, to train a model, we convert the movie data into a sparse matrix of 0's and 1's, where each column is a movie genre, and each row is a movie title. This matrix is `genre_mat2` and we can see its truncated version in the output below (though `str()` transposes the matrix visually, flipping movie titles/genres):

```
movie_genre <- as.data.frame(movie_data$genres, stringsAsFactors = FALSE)
movie_genre2 <- as.data.frame(
  tstrsplit(movie_genre[, 1], "[|]",
    type.convert = TRUE
  ),
  stringsAsFactors = FALSE
)
colnames(movie_genre2) <- c(1:10)

list_genre <- c(
  "Action", "Adventure", "Animation", "Children",
  "Comedy", "Crime", "Documentary", "Drama", "Fantasy",
  "Film-Noir", "Horror", "Musical", "Mystery", "Romance",
  "Sci-Fi", "Thriller", "War", "Western"
)
genre_mat1 <- matrix(0, 10330, 18)
genre_mat1[1, ] <- list_genre
```

```

colnames(genre_mat1) <- list_genre

for (index in 1:nrow(movie_genre2)) {
  for (col in 1:ncol(movie_genre2)) {
    gen_col <- which(genre_mat1[1, ] == movie_genre2[index, col])
    genre_mat1[index + 1, gen_col] <- 1
  }
}

# remove first row, which was the genre list
genre_mat2 <- as.data.frame(genre_mat1[-1, ], stringsAsFactors = FALSE)
for (col in 1:ncol(genre_mat2)) {
  genre_mat2[, col] <- as.integer(genre_mat2[, col]) # convert from characters to integers
}
str(genre_mat2)

```

```

## 'data.frame': 10329 obs. of 18 variables:
## $ Action : int 0 0 0 0 0 1 0 0 1 1 ...
## $ Adventure : int 1 1 0 0 0 0 0 1 0 1 ...
## $ Animation : int 1 0 0 0 0 0 0 0 0 0 ...
## $ Children : int 1 1 0 0 0 0 0 1 0 0 ...
## $ Comedy : int 1 0 1 1 1 0 1 0 0 0 ...
## $ Crime : int 0 0 0 0 0 1 0 0 0 0 ...
## $ Documentary: int 0 0 0 0 0 0 0 0 0 0 ...
## $ Drama : int 0 0 0 1 0 0 0 0 0 0 ...
## $ Fantasy : int 1 1 0 0 0 0 0 0 0 0 ...
## $ Film-Noir : int 0 0 0 0 0 0 0 0 0 0 ...
## $ Horror : int 0 0 0 0 0 0 0 0 0 0 ...
## $ Musical : int 0 0 0 0 0 0 0 0 0 0 ...
## $ Mystery : int 0 0 0 0 0 0 0 0 0 0 ...
## $ Romance : int 0 0 1 1 0 0 1 0 0 0 ...
## $ Sci-Fi : int 0 0 0 0 0 0 0 0 0 0 ...
## $ Thriller : int 0 0 0 0 0 1 0 0 0 1 ...
## $ War : int 0 0 0 0 0 0 0 0 0 0 ...
## $ Western : int 0 0 0 0 0 0 0 0 0 0 ...

```

Next, we create a search matrix that merge movie_data and genre_mat2. This allows user to easily perform a search of the movie titles based on the genres listed in our list. This merged matrix is SearchMatrix below.

```

# Using column bind function to merge movie_data and genre_mat2 dataframes
SearchMatrix <- cbind(movie_data[, 1:2], genre_mat2[])
head(SearchMatrix)

```

```

##   movieId      title Action Adventure Animation
## 1      1      Toy Story (1995)      0      1      1
## 2      2      Jumanji (1995)      0      1      0
## 3      3      Grumpier Old Men (1995)      0      0      0
## 4      4      Waiting to Exhale (1995)      0      0      0
## 5      5      Father of the Bride Part II (1995)      0      0      0
## 6      6      Heat (1995)      1      0      0
##   Children Comedy Crime Documentary Drama Fantasy Film-Noir Horror Musical
## 1      1      1      0      0      0      1      0      0      0
## 2      1      0      0      0      0      1      0      0      0
## 3      0      1      0      0      0      0      0      0      0
## 4      0      1      0      0      1      0      0      0      0
## 5      0      1      0      0      0      0      0      0      0

```

```
## 6      0      0      1      0      0      0      0      0      0
##  Mystery Romance Sci-Fi Thriller War Western
## 1      0      0      0      0      0      0
## 2      0      0      0      0      0      0
## 3      0      1      0      0      0      0
## 4      0      1      0      0      0      0
## 5      0      0      0      0      0      0
## 6      0      0      0      1      0      0
```

Then, we create a sparse matrix of ratings. Each row represents a user and each column is a movie. That makes each entry in the matrix (`ratingMatrix`) a rating given by a particular user for a particular movie. The matrix is sparse because most movies are not rated by most users. The output below shows the matrix dimensions. We force that object to be of `realRatingMatrix` type, for use by our `recommenderlab` Machine Learning package later.

```
ratingMatrix <- dcast(
  rating_data, userId ~ movieId,
  value.var = "rating", na.rm = FALSE
)
ratingMatrix <- as.matrix(ratingMatrix[, -1]) # remove userIds
# Convert rating matrix into a recommenderlab sparse matrix
ratingMatrix <- as(ratingMatrix, "realRatingMatrix")
ratingMatrix
```

```
## 668 x 10325 rating matrix of class 'realRatingMatrix' with 105339 ratings.
```

We now filter out data to only keep what is useful. Let us only keep that have “enough” data. “Enough” in this case is films rated by more than 50 users and users who have rated more than 50 films.

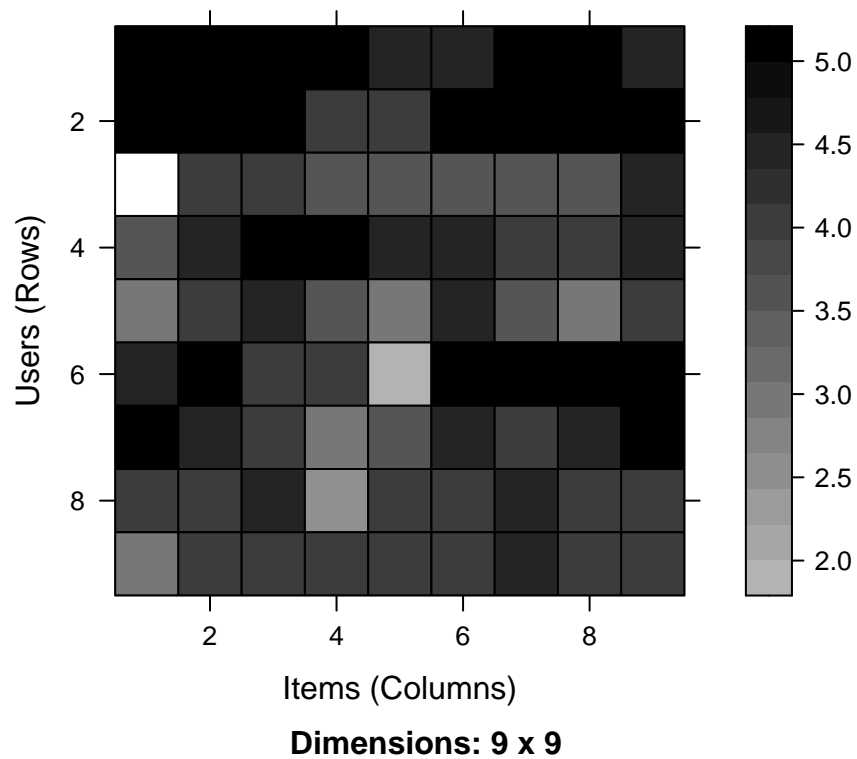
```
movie_ratings <- ratingMatrix[
  rowCounts(ratingMatrix) > 50,
  colCounts(ratingMatrix) > 50
]
movie_ratings
```

```
## 420 x 447 rating matrix of class 'realRatingMatrix' with 38341 ratings.
```

We can visualize top users/top movies with this heatmap; the darker the square, the higher the rating given by that particular user (indicated by a row) given to that particular movie (given by a column):

```
minimum_movies <- quantile(rowCounts(movie_ratings), 0.98)
minimum_users <- quantile(colCounts(movie_ratings), 0.98)
image(
  movie_ratings[
    rowCounts(movie_ratings) > minimum_movies,
    colCounts(movie_ratings) > minimum_users
  ],
  main = "Heatmap of the top users and movies"
)
```

Heatmap of the top users and movies



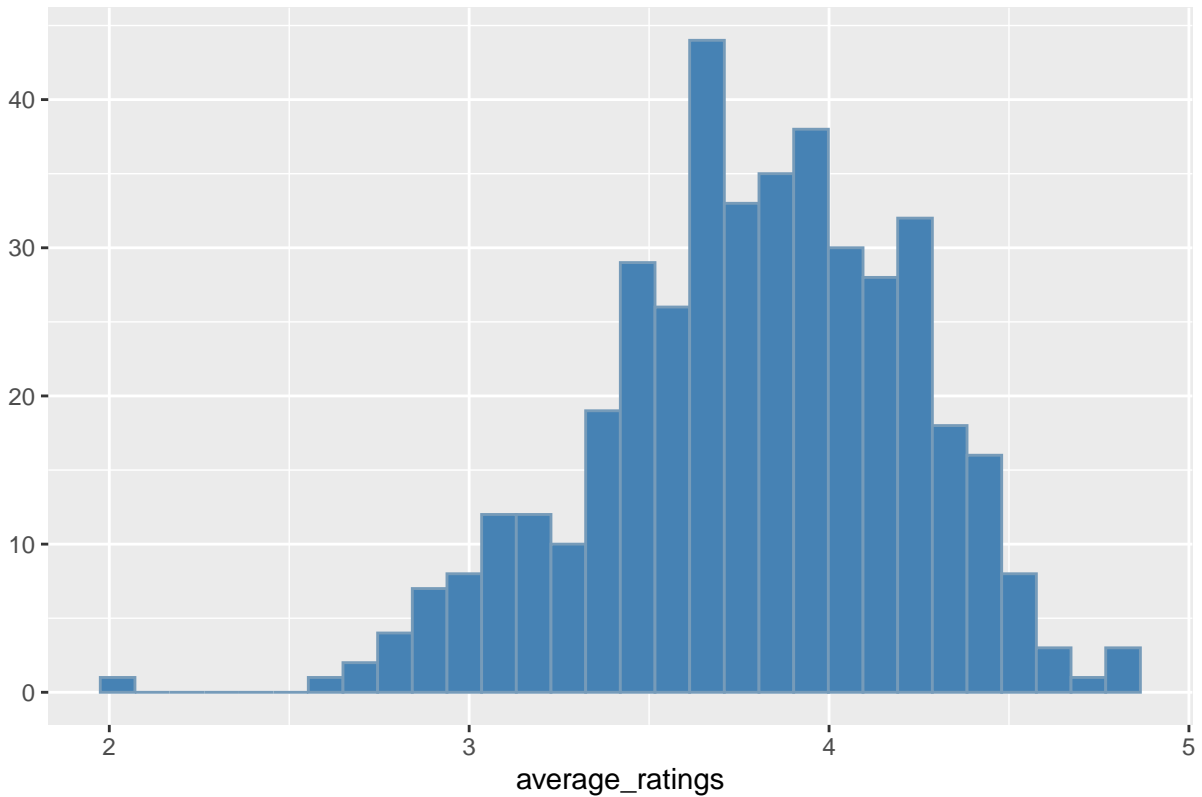
Let us visualize distribution of the average ratings per user:

```
average_ratings <- rowMeans(movie_ratings)
qplot(
  average_ratings,
  fill = I("steelblue"),
  col = I("#769bb9")
) +
  ggtitle("Distribution of the average rating per user")
```

Warning: `qplot()` was deprecated in ggplot2 3.4.0.

`stat_bin()` using `bins = 30`. Pick better value with `binwidth`.

Distribution of the average rating per user



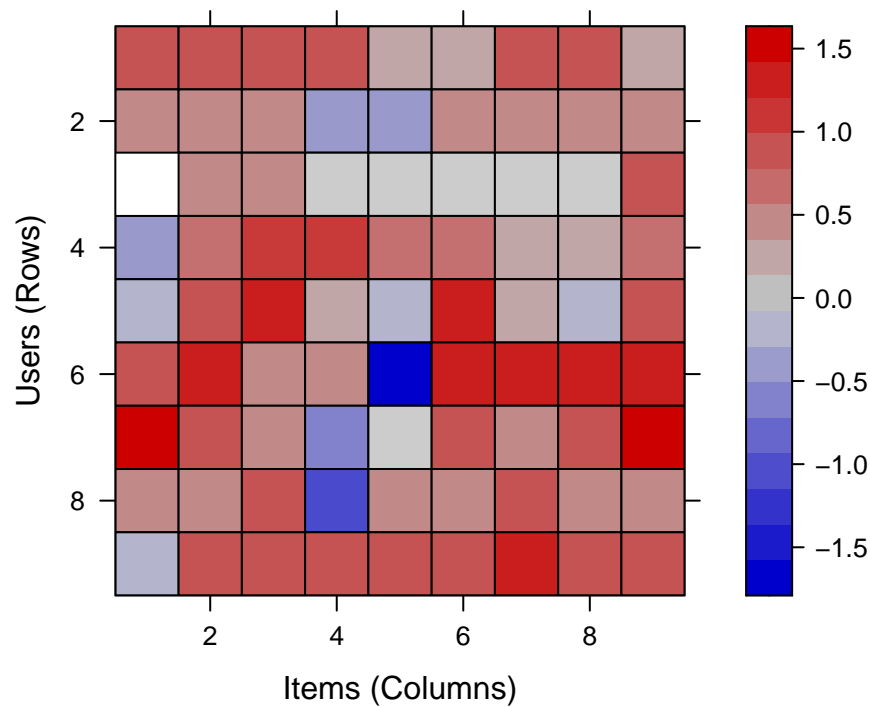
The next step in data preparation is to normalize user ratings to have mean of 0 to limit bias of extreme ratings on the model training later:

```
normalized_ratings <- normalize(movie_ratings)
sum(rowMeans(normalized_ratings) > 0.00001)
```

```
## [1] 0
```

```
image(
  normalized_ratings[
    rowCounts(normalized_ratings) > minimum_movies,
    colCounts(normalized_ratings) > minimum_users
  ],
  main = "Normalized Ratings of the Top Users"
)
```

Normalized Ratings of the Top Users



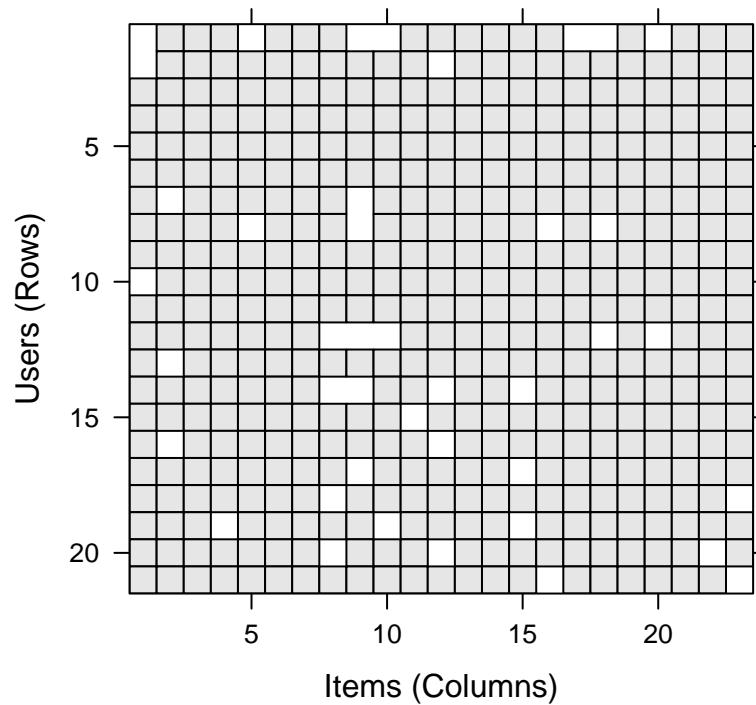
Dimensions: 9 x 9

As our final step in data preparation, we will binarize this data to have discrete values of 0 and 1. This allows us to create a matrix that will consist of 1 if the movie rating is above 3 and 0 otherwise. This will allow for a more efficient recommendation system later; this is also visualized:

```
binary_minimum_movies <- quantile(rowCounts(movie_ratings), 0.95)
binary_minimum_users <- quantile(colCounts(movie_ratings), 0.95)

goodRatedFilms <- binarize(movie_ratings, minRating = 3)
image(
  goodRatedFilms[
    rowCounts(movie_ratings) > binary_minimum_movies,
    colCounts(movie_ratings) > binary_minimum_users
  ],
  main = "Heatmap of the top users and movies"
)
```

Heatmap of the top users and movies



Dimensions: 21 x 23

6. Discuss the modeling. What modeling was used?

We will first split our dataset into 80% training set `training_data` and 20% test set `testing_data`.

```
# splitting the dataset to 80% training set and 20% test set
set.seed(1234) # to generate reproducible results
sampled_data <- sample(
  x = c(TRUE, FALSE),
  size = nrow(movie_ratings),
  replace = TRUE,
  prob = c(0.8, 0.2)
)
training_data <- movie_ratings[sampled_data, ]
testing_data <- movie_ratings[!sampled_data, ]
```

For the modeling part of this project, we will be utilizing Item Based Collaborative Filtering System. This type of collaborative filtering looks for similarities between items based on the movie ratings. Since the parameters are default by nature, we will set `k` parameter to 30. `k` represents the number of items for computing similarities, the algorithm will identify `k` most similar items and store their respective number.

```
recommen_model <- Recommender(
  data = training_data,
  method = "IBCF",
  parameter = list(k = 30)
)
recommen_model
```



```
## Recommender of type 'IBCF' for 'realRatingMatrix'
## learned using 342 users.
```

```
class(recommen_model)
```

```
## [1] "Recommender"
## attr("package")
## [1] "recommenderlab"
```

After we retrieve the `recommen_model`, we then generate a heatmap that contains top 20 films from our training set.

```
model_info <- getModel(recommen_model)
class(model_info$sim)
```

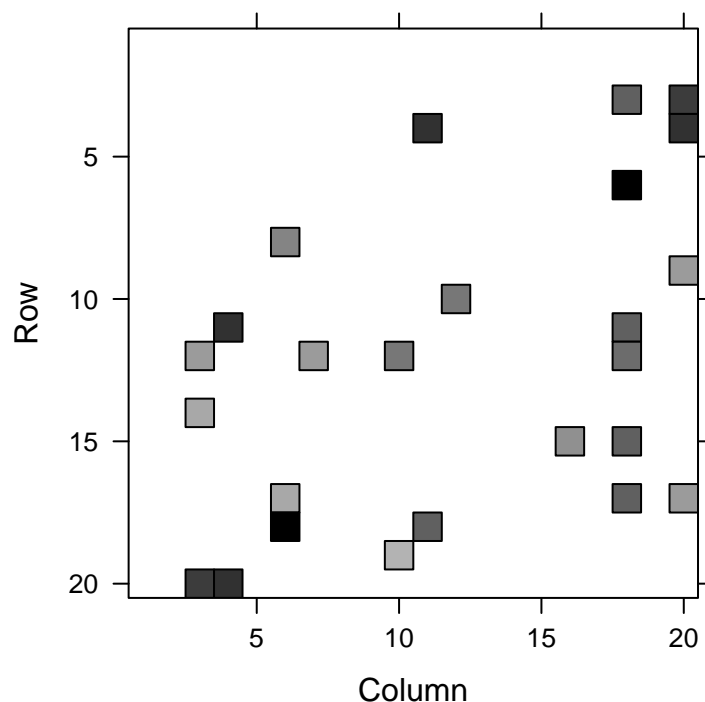
```
## [1] "dgCMatrix"
## attr("package")
## [1] "Matrix"
```

```
dim(model_info$sim)
```

```
## [1] 447 447
```

```
top_items <- 20
image(model_info$sim[1:top_items, 1:top_items],
      main = "Heatmap of the first rows and columns")
```

Heatmap of the first rows and columns

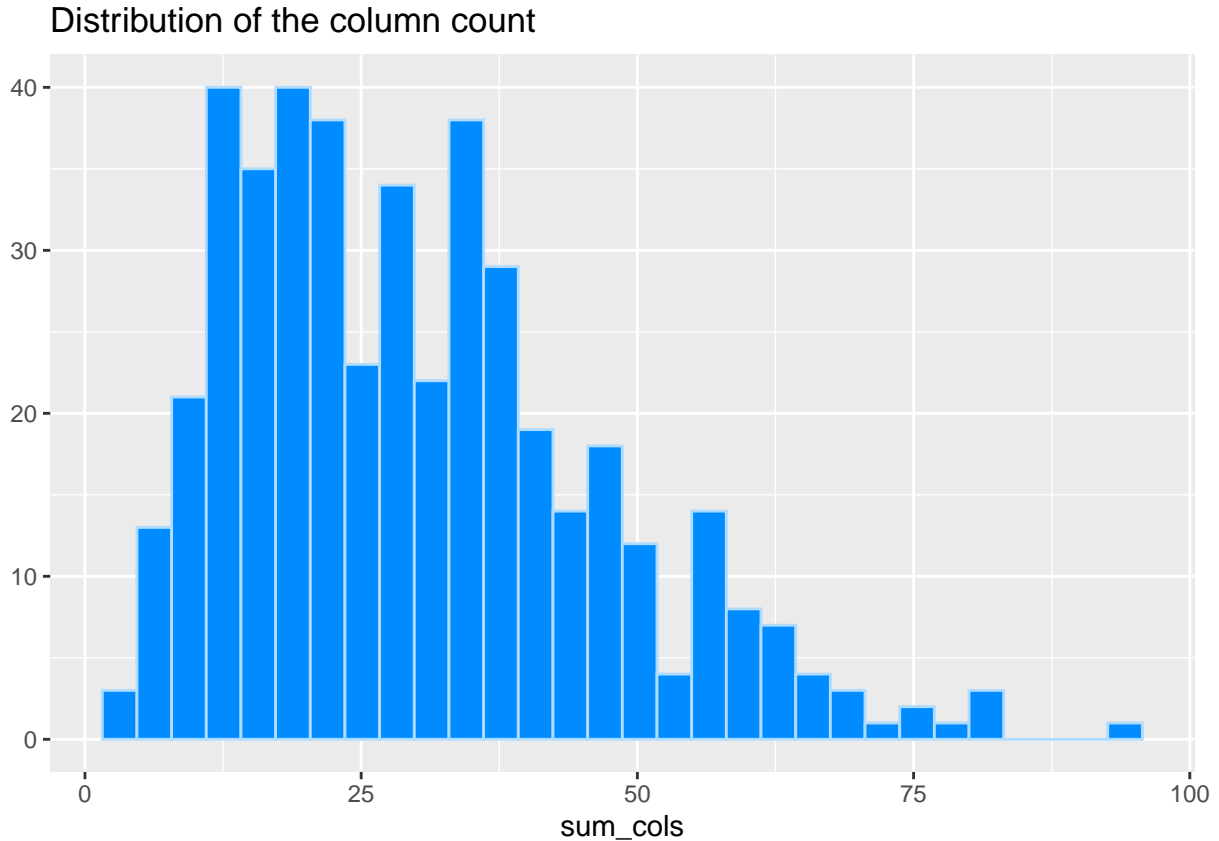


Dimensions: 20 x 20

Let us visualize the distribution of the sums of similarities per each movie. Which similarity score sums are the most frequent?

```
sum_cols <- colSums(model_info$sim > 0)
qplot(sum_cols, fill = I("#008cff"), col = I("#afdbff")) +
  ggtitle("Distribution of the column count")
```

`stat_bin()` using `bins = 30`. Pick better value with `binwidth`.



7. Produce and discuss the output. What are the results?

By using `predict()` function, we can identify similar films and rank them. For the `top_recommendations` variable, we are setting it to 10 which is the number of films that will be recommended to each user. Note that we will be using the testing set here.

```
top_recommendations <- 10 # the number of items to recommend to each user
predicted_recommendations <- predict(object = recommen_model,
                                     newdata = testing_data,
                                     n = top_recommendations)
predicted_recommendations
```

Recommendations as 'topNList' with n = 10 for 78 users.

Below is an example of what the `predicted_recommendations` function will do. By using the recommendation given for the first user, we can use that to recommend movies for the second user.

```
# recommendation for the first user
user1 <- predicted_recommendations@items[[1]]
movies_user1 <- predicted_recommendations@itemLabels[user1]
movies_user2 <- movies_user1
for (index in 1:10){
```

```

    movies_user2[index] <- as.character(
      subset(movie_data, movie_data$movieId == movies_user1[index])$title
    )
  }
  movies_user2

```

```

## [1] "Braveheart (1995)"
## [2] "Rumble in the Bronx (Hont faan kui) (1995)"
## [3] "Congo (1995)"
## [4] "Judge Dredd (1995)"
## [5] "Outbreak (1995)"
## [6] "Pulp Fiction (1994)"
## [7] "Shawshank Redemption, The (1994)"
## [8] "Star Trek: Generations (1994)"
## [9] "Flintstones, The (1994)"
## [10] "Forrest Gump (1994)"

```

recommendation_matrix has 10 rows (i.e. 10 movie recommendations per user) and each column represents a user (by user ID). Each entry is an ID of a recommended movie.

```

# matrix with the recommendations for each user
recommendation_matrix <- sapply(predicted_recommendations@items,
                                function(x){ as.integer(colnames(movie_ratings)[x]) })
#dim(recc_matrix)
recommendation_matrix[,1:4]

```

```

##           0      1      2      3
## [1,] 110  168  235   16
## [2,] 112  508 2005   17
## [3,] 160  661 2167   34
## [4,] 173  858 3703  539
## [5,] 292 1047  924  969
## [6,] 296 1080  858 1201
## [7,] 318 1302 1036 1240
## [8,] 329 1580 5060 1249
## [9,] 355 1961 1278 1358
## [10,] 356 2353 48516 2542

```

8. Provide an explanation with any visuals.

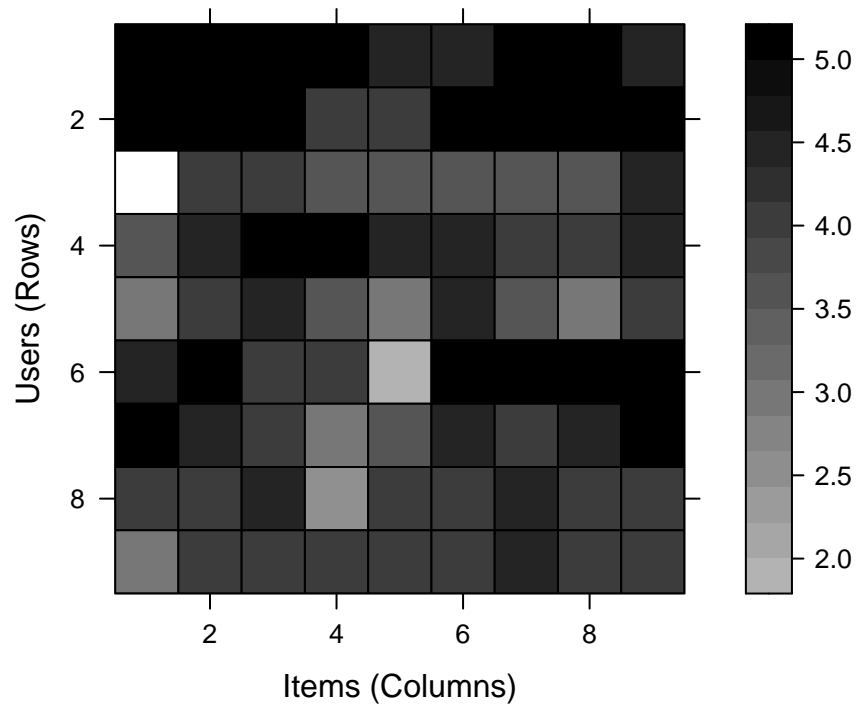
Previously in (5), we've done visualizations with heatmaps. The first one was top users/top movies heatmap; the darker the square, the higher the rating given by that particular user (indicated by a row) given to that particular movie (given by a column). Here's a reproduction of it from earlier:

```

image(
  movie_ratings[
    rowCounts(movie_ratings) > minimum_movies,
    colCounts(movie_ratings) > minimum_users
  ],
  main = "Heatmap of the top users and movies"
)

```

Heatmap of the top users and movies

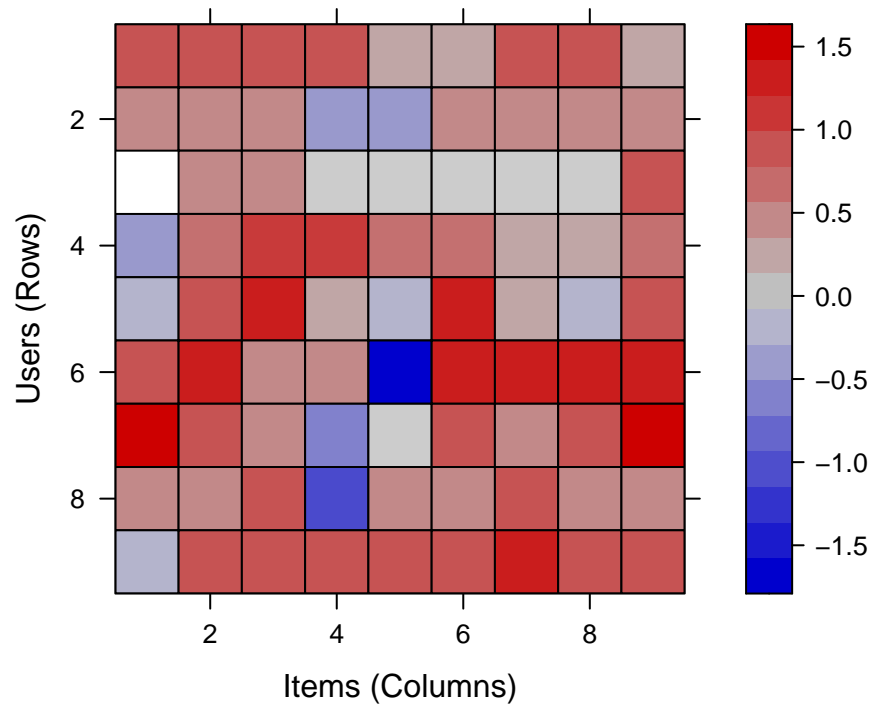


Dimensions: 9 x 9

Similarly, here is a reproduction of the normalized version of these ratings from (5):

```
image(  
  normalized_ratings[  
    rowCounts(normalized_ratings) > minimum_movies,  
    colCounts(normalized_ratings) > minimum_users  
  ],  
  main = "Normalized Ratings of the Top Users"  
)
```

Normalized Ratings of the Top Users

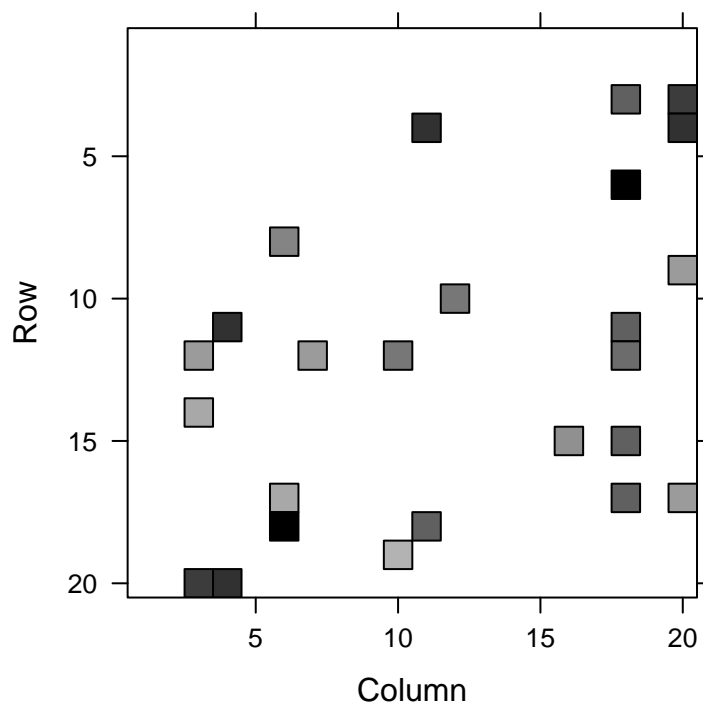


Dimensions: 9 x 9

After training our recommendation, we generated a heatmap that contains top 20 films from our training set; to visualize, among top 20, which ones are most similar to which other movies. This is exploring what will be the basis of our recommendations (recommending movies similar to what a user already liked).

```
image(model_info$sim[1:top_items, 1:top_items],  
      main = "Heatmap of the first rows and columns")
```

Heatmap of the first rows and columns



Dimensions: 20 x 20

From the `recommendation_matrix` that we generated above, we can visualize the distribution (by count) of the top 10 recommended movie titles via this histogram.

```
recommendation_table <- as.data.frame(
  as.vector(recommendation_matrix)
)
colnames(recommendation_table)[1] <- "movieId"

recommendation_table_top <- recommendation_table %>%
  group_by(movieId) %>%
  filter(n() > 5)

recommendation_table_top$title <- with(
  movie_data, title[match(recommendation_table_top$movieId, movieId)]
)

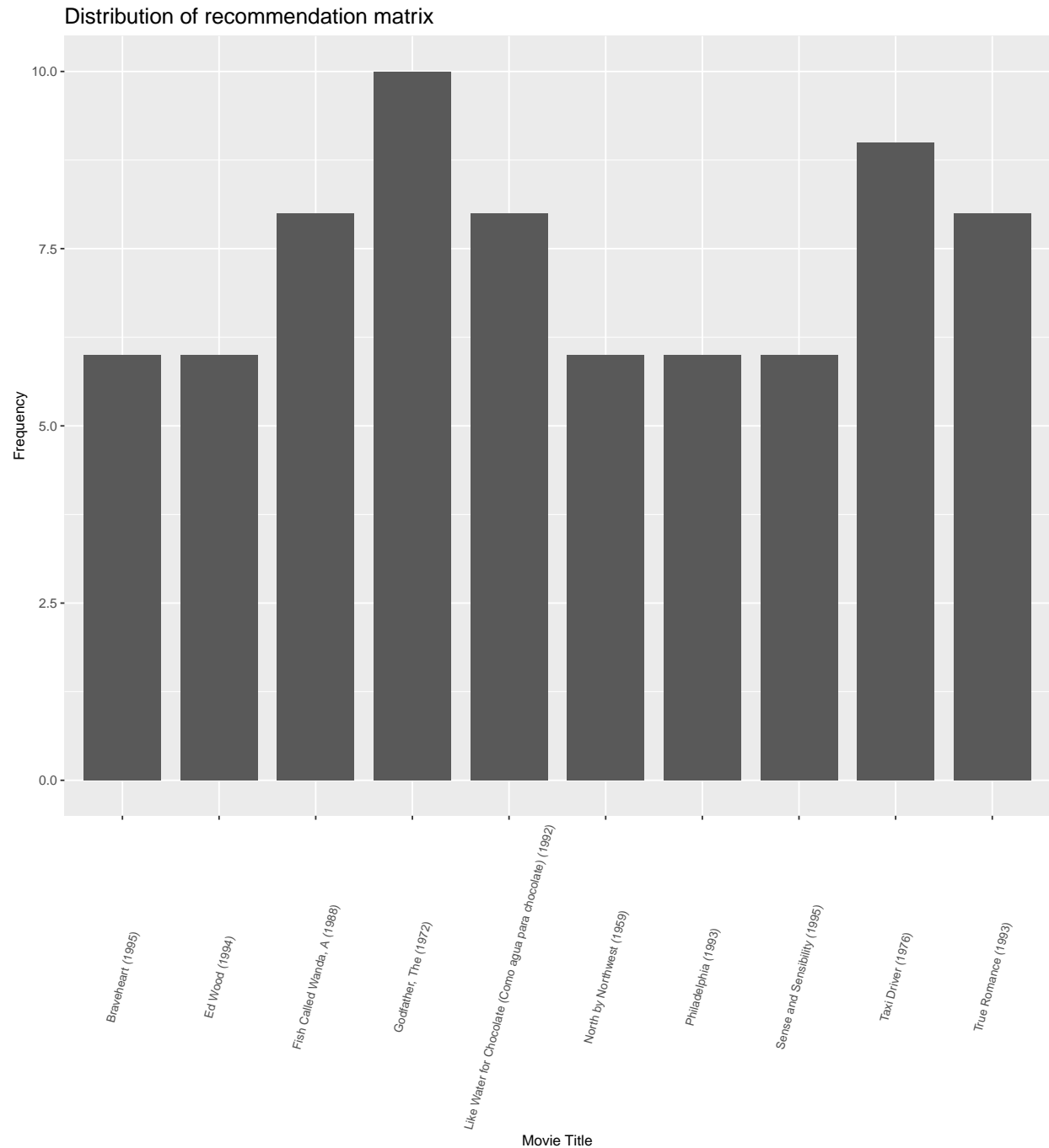
recommendation_table_top <- recommendation_table_top %>%
  add_count(title, sort = TRUE) %>%
  distinct()

ggplot(recommendation_table_top[1:10, ], aes(x = title, y = n)) +
  geom_bar(stat = "identity", show.legend = FALSE, width = 0.8) +
  labs(x = "Movie Title",
       y = "Frequency",
       title = "Distribution of recommendation matrix") +
  theme(axis.text.x = element_text(
    angle = 75, vjust = 0.5, hjust = 0.5, size = 8
  ))
```

```

),
axis.title.x = element_text(size = 10),
axis.title.y = element_text(size = 10),
plot.title = element_text(size = 15)
)

```



So we are seeing the most frequently recommended movie is Godfather, The (1972).

Citations

DataFlair. 2023. “Machine Learning Project – Data Science Movie Recommendation System Project in R.”

DataFlair. <https://data-flair.training/blogs/data-science-r-movie-recommendation/>.