

2 Hands On: Web Usage Mining

Read the file `log.csv`, containing information on the web pages visited by a set of users, into a data frame in R.

```
library(dplyr)
library(readr)
d <- read_csv("log.csv")
```

2.1 Simple Recommendation Strategies

Most Visited Pages

1. Recommend the 3 most visited pages.

For that purpose:

- (a) inspect how many times each page was visited;

```
d %>% group_by(PAGE) %>% tally()

## # A tibble: 7 x 2
##   PAGE     n
##   <chr> <int>
## 1 A         3
## 2 B         3
## 3 C         4
## 4 F         2
## 5 G         2
## 6 I         2
## 7 J         1
```

- (b) sort the pages by decreasing number of visits;

```
d %>% group_by(PAGE) %>% tally(sort=TRUE)

## # A tibble: 7 x 2
##   PAGE     n
##   <chr> <int>
## 1 C         4
## 2 A         3
## 3 B         3
## 4 F         2
## 5 G         2
## 6 I         2
## 7 J         1
```

- (c) obtain the top 3 pages for recommendation.

```
d %>% group_by(PAGE) %>% tally(sort=TRUE) %>% top_n(3) %>% pull(PAGE)

## [1] "C" "A" "B"
```

Using Clustering Results

2. Suppose we want to form two clusters of users, according to the pages they have visited.

For that purpose:

- (a) start by transforming the log access data into a matrix that has on each row a user and for each user the information on his visits to each page; this can be obtained with the `table()` function;

```
dat <-table(d$USER,d$PAGE)
```

```
##
##      A B C F G I J
## u1 1 1 1 0 0 0 0
## u2 1 0 1 0 0 0 0
## u3 0 1 0 1 1 1 0
## u4 0 1 1 0 0 0 0
## u5 0 0 0 1 1 1 1
## u6 1 0 1 0 0 0 0
```

- (b) use the function `dist()` to obtain a distance matrix with the Euclidean distance between the users;

```
library(proxy)
dm <-dist(dat) # euclidean
summary(pr_DB)
```

```
## * Similarity measures:
## Braun-Blanquet, Chi-squared, correlation, cosine, Cramer, Dice, eDice,
## eJaccard, Fager, Faith, Gower, Hamman, Jaccard, Kulczynski1,
## Kulczynski2, Michael, Mountford, Mozley, Ochiai, Pearson, Phi,
## Phi-squared, Russel, simple matching, Simpson, Stiles, Tanimoto,
## Tschuprow, Yule, Yule2
##
## * Distance measures:
## Bhjattacharyya, Bray, Canberra, Chord, divergence, Euclidean, fJaccard,
## Geodesic, Hellinger, Kullback, Levenshtein, Mahalanobis, Manhattan,
## Minkowski, Podani, Soergel, supremum, Wave, Whittaker
```

```
dm <- dist(dat,method="jaccard")
```

```
##      u1      u2      u3      u4      u5
## u2 0.333333
## u3 0.833333 1.000000
## u4 0.333333 0.666667 0.800000
## u5 1.000000 1.000000 0.400000 1.000000
## u6 0.333333 0.000000 1.000000 0.666667 1.000000
```

- (c) check for alternatives in the help page of `dist()`;
- (d) use the function `hclust()` with the distance matrix to obtain an agglomerative clustering model of this data;

```
c1 <- hclust(dm)
```

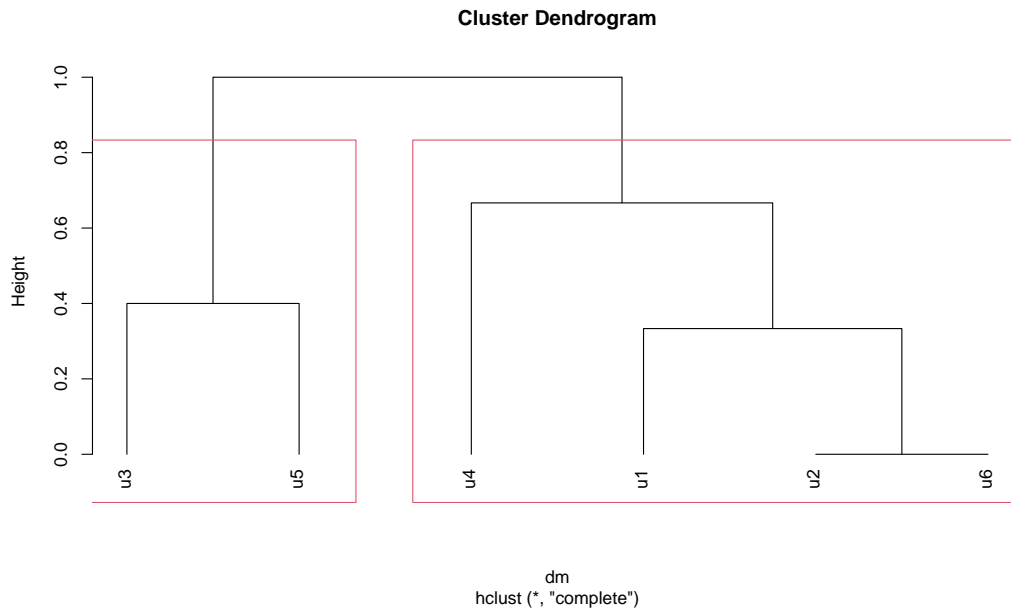
- (e) visualize the obtained dendrogram with function `plot()`;
- (f) visualize again the dendrogram, but now with option `hang=-0.1`.
- (g) use the function `cutree()` to "cut" the hierarchical clustering in just two clusters; inspect the cluster membership of each user;

```
cm <- cutree(c1,2)
```

```
## u1 u2 u3 u4 u5 u6
## 1 1 2 1 2 1
```

(h) use the function `rect.hclust()` to draw the previous solution in the dendrogram.

```
plot(c1, hang=-0.1)
rect.hclust(c1, 2)
```



3. Recommend the top 2 pages for users of cluster 1.

For that purpose:

(a) inspect what were the pages visited by users in cluster 1;

```
dd <- mutate(d, Cluster=cm[d$USER]) # adding the cluster of each user
dd
```

```
## # A tibble: 17 x 3
##   USER PAGE Cluster
##   <chr> <chr> <int>
## 1 u1    A      1
## 2 u1    B      1
## 3 u1    C      1
## 4 u2    A      1
## 5 u2    C      1
## 6 u3    B      2
## 7 u3    G      2
## 8 u3    F      2
## 9 u3    I      2
## 10 u4   B      1
## 11 u4   C      1
## 12 u5   G      2
## 13 u5   F      2
## 14 u5   I      2
## 15 u5   J      2
## 16 u6   A      1
## 17 u6   C      1
```

```
filter(dd, Cluster==1) %>% pull(PAGE) # the answer (the pages)
```

```
## [1] "A" "B" "C" "A" "C" "B" "C" "A" "C"
```

(b) inspect how many times each of these pages were visited;

```
filter(dd,Cluster==1) %>% group_by(PAGE) %>% tally()

## # A tibble: 3 x 2
##   PAGE     n
##   <chr> <int>
## 1 A         3
## 2 B         2
## 3 C         4
```

(c) sort the pages by decreasing order of visits;

```
filter(dd,Cluster==1) %>% group_by(PAGE) %>% tally(sort=TRUE)

## # A tibble: 3 x 2
##   PAGE     n
##   <chr> <int>
## 1 C         4
## 2 A         3
## 3 B         2
```

(d) obtain the top 2 pages for recommendation.

```
filter(dd,Cluster==1) %>%
  group_by(PAGE) %>% tally(sort=TRUE) %>% top_n(2) %>% pull(PAGE)

## [1] "C" "A"
```

4. Recommend the top 2 pages for users of cluster 2.

```
filter(dd,Cluster==1) %>%
  group_by(PAGE) %>% tally(sort=TRUE) %>% top_n(2) %>% pull(PAGE)

## [1] "C" "A"
```

5. Using the same clustering results, recommend the top 3 pages for user u2.

From that top pages you should remove the pages that the user has already visited.

```
cluster.u2 <- dd %>% filter(USER == "u2") %>% select(Cluster) %>% head(1) %>% pull()

rec.u2 <- filter(dd,Cluster==cluster.u2) %>%
  group_by(PAGE) %>% tally(sort=TRUE) %>% top_n(3) %>% select(PAGE)

seen.u2 <- dd %>% filter(USER == "u2") %>% select(PAGE)

anti_join(rec.u2,seen.u2)

## # A tibble: 1 x 1
##   PAGE
##   <chr>
## 1 B
```

Load the package recommenderlab.

2.2 Recommendation using Association Rules

6. Obtain a recommendation model using association rules with the first 6 users. For that purpose:

(a) start by coercing the data frame with user-page access information from the log1.csv file to a binaryRatingMatrix (brm);

```
log <- read_csv("log1.csv", col_types = list(col_factor(), col_factor()))
brm <- as(as.data.frame(log), "binaryRatingMatrix")
```

- (b) select the information on the first 6 users to be used as training offline data and save it to a new variable (e.g brm_offline);

```
brm_offline <- brm[1:6,]
```

- (c) inspect the content of brm_offline; use the function getRatingMatrix and getData.frame;

```
getData.frame(brm_offline)

##      user item rating
## 1    u1    A      1
## 4    u1    B      1
## 7    u1    C      1
## 2    u2    A      1
## 8    u2    C      1
## 5    u3    B      1
## 11   u3    G      1
## 13   u3    F      1
## 15   u3    I      1
## 6    u4    B      1
## 9    u4    C      1
## 12   u5    G      1
## 14   u5    F      1
## 16   u5    I      1
## 17   u5    J      1
## 3    u6    A      1
## 10   u6    C      1

getRatingMatrix(brm_offline)

## itemMatrix in sparse format with
## 6 rows (elements/transactions) and
## 7 columns (items)

inspect(getRatingMatrix(brm_offline))

##      items
## [1] {A,B,C}
## [2] {A,C}
## [3] {B,G,F,I}
## [4] {B,C}
## [5] {G,F,I,J}
## [6] {A,C}
```

- (d) apply the functions rowCounts and colCounts to brm_offline; what information does it give you?

```
rowCounts(brm_offline)

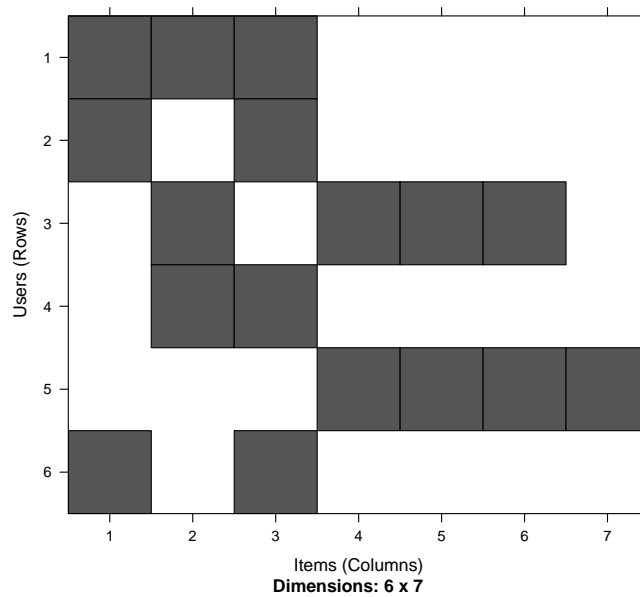
## u1 u2 u3 u4 u5 u6
## 3 2 4 2 4 2

colCounts(brm_offline)

## A B C G F I J
## 3 3 4 2 2 2 1
```

- (e) apply the function image to brm_offline;

```
image(brm_offline)
```



- (f) obtain the recommender model based on association rules with the instruction
- ```
modelAR <- Recommender(brm_offline,"AR")
```

```
model <- Recommender(brm_offline, "AR")
```

- (g) apply the function `getModel` to the obtained model and then inspect the association rules that compose the model.

```
getModel(model)

$description
[1] "AR: rule base"
##
$rule_base
set of 26 rules
##
$support
[1] 0.1
##
$confidence
[1] 0.8
##
$maxlen
[1] 3
##
$sort_measure
[1] "confidence"
##
$sort_decreasing
[1] TRUE
##
$apriori_control
$apriori_control$verbose
[1] FALSE
##
$verbose
[1] FALSE

rules <- getModel(model)$rule_base
inspect(rules)
```

```
lhs rhs support confidence lift count
[1] {J} => {G} 0.1666667 1 3.0 1
[2] {J} => {F} 0.1666667 1 3.0 1
[3] {J} => {I} 0.1666667 1 3.0 1
[4] {A} => {C} 0.5000000 1 1.5 3
[5] {G} => {F} 0.3333333 1 3.0 2
[6] {F} => {G} 0.3333333 1 3.0 2
[7] {G} => {I} 0.3333333 1 3.0 2
[8] {I} => {G} 0.3333333 1 3.0 2
[9] {F} => {I} 0.3333333 1 3.0 2
[10] {I} => {F} 0.3333333 1 3.0 2
[11] {G,J} => {F} 0.1666667 1 3.0 1
[12] {F,J} => {G} 0.1666667 1 3.0 1
[13] {G,J} => {I} 0.1666667 1 3.0 1
[14] {I,J} => {G} 0.1666667 1 3.0 1
[15] {F,J} => {I} 0.1666667 1 3.0 1
[16] {I,J} => {F} 0.1666667 1 3.0 1
[17] {A,B} => {C} 0.1666667 1 1.5 1
[18] {G,F} => {I} 0.3333333 1 3.0 2
[19] {G,I} => {F} 0.3333333 1 3.0 2
[20] {F,I} => {G} 0.3333333 1 3.0 2
[21] {B,G} => {F} 0.1666667 1 3.0 1
[22] {B,F} => {G} 0.1666667 1 3.0 1
[23] {B,G} => {I} 0.1666667 1 3.0 1
[24] {B,I} => {G} 0.1666667 1 3.0 1
[25] {B,F} => {I} 0.1666667 1 3.0 1
[26] {B,I} => {F} 0.1666667 1 3.0 1
```

7. Suppose that u7 enters the system and becomes an active user. Deploy the recommendation model for him/her.

For that purpose:

- apply the predict function with the model and the rating matrix of the user, such that only the top 2 recommendations are given as output;
- apply the function getList to the obtained predictions to inspect the actual recommendations; which are they?
- to comprove the obtained recommendations, filter the rules which have been triggered for this active user.

```
brm_u7 <- brm[7,]
recsAR <- predict(model, brm_u7, n=2)
recsAR

Recommendations as 'topNList' with n = 2 for 1 users.

getList(recsAR)

$u7
[1] "G" "I"

r <- subset(rules,lhs %in% c("C","F"))
inspect(r)

lhs rhs support confidence lift count
[1] {F} => {G} 0.3333333 1 3 2
[2] {F} => {I} 0.3333333 1 3 2
[3] {F,J} => {G} 0.1666667 1 3 1
[4] {F,J} => {I} 0.1666667 1 3 1
[5] {G,F} => {I} 0.3333333 1 3 2
[6] {F,I} => {G} 0.3333333 1 3 2
[7] {B,F} => {G} 0.1666667 1 3 1
[8] {B,F} => {I} 0.1666667 1 3 1
```

8. Now suppose that u8 enters the system and becomes an active user. Deploy the recommendation model for him/her. Be critical regarding the results.

```
brm_u8 <- brm[8,]
recsAR <- predict(model, brm_u8, n=2)
recsAR

Recommendations as 'topNList' with n = 2 for 1 users.

getList(recsAR)

$u8
character(0)

r <- subset(rules,lhs %in% c("C"))
inspect(r)
```

9. Explore the types of recommendation models available for binary rating matrices.

```
recommenderRegistry$get_entries(dataType ="binaryRatingMatrix")
```

10. Make the top 2 recommendations to u7 and u8 using the popularity of the pages, instead of association rules. Try to understand the obtained recommendations.

```
modelPop <- Recommender(brm_offline,"POPULAR")
recsPop <- predict(modelPop, brm[7:8,], n=2)
recsPop

Recommendations as 'topNList' with n = 2 for 2 users.

getList(recsPop)

$u7
[1] "B" "A"
##
$u8
[1] "B" "A"
```

## 2.3 Recommendation using Collaborative Filtering

### Binary Rating Data

Considering the same binary rating matrix of the previous exercise `brm_offline`, build a recommendation model based on collaborative filtering.

11. Start by using the function `similarity` to build the similarity cosine matrix for:

(a) an user-based approach;

```
simCos_users<- similarity(brm_offline,method="cosine")
simCos_users

u1 u2 u3 u4 u5
u2 0.8164966
u3 0.2886751 0.0000000
u4 0.8164966 0.5000000 0.3535534
u5 0.0000000 0.0000000 0.7500000 0.0000000
u6 0.8164966 1.0000000 0.0000000 0.5000000 0.0000000
```

(b) an item-based approach.

```
simCos_items <- similarity(brm_offline,method="cosine",which="items")
simCos_items

A B C G F I
B 0.3333333
C 0.8660254 0.5773503
G 0.0000000 0.4082483 0.0000000
F 0.0000000 0.4082483 0.0000000 1.0000000
I 0.0000000 0.4082483 0.0000000 1.0000000 1.0000000
J 0.0000000 0.0000000 0.0000000 0.7071068 0.7071068 0.7071068
```



12. Obtain the top 2 recommendations with user-based CF and item-based CF methods using the cosine similarity with a neighborhood of size 3, for:

```
modelUBCF <- Recommender(brm_offline, "UBCF",parameter=list(method="cosine",nn=3)) #weighted=FALSE
getModel(modelUBCF)

$description
[1] "UBCF-Binary Data: contains full or sample of data set"
##
$data
6 x 7 rating matrix of class 'binaryRatingMatrix' with 17 ratings.
##
$method
[1] "cosine"
##
$nn
[1] 3
##
$weighted
[1] TRUE
##
$sample
[1] FALSE
##
$verbose
[1] FALSE

modelIBCF <- Recommender(brm_offline, "IBCF",parameter=list(method="cosine",k=3))
getModel(modelIBCF)

$description
[1] "IBCF: Reduced similarity matrix"
##
$sim
7 x 7 sparse Matrix of class "dgCMatrix"
A B C G F I J
A . 0.3333333 0.8660254
B . . 0.5773503 . 0.4082483 0.4082483 .
C 0.8660254 0.5773503
G . . . 1.0000000 1.0000000 0.7071068
F . . . 1.0000000 . 1.0000000 0.7071068
I . . . 1.0000000 1.0000000 . 0.7071068
J . . . 0.7071068 0.7071068 0.7071068 .
##
$k
[1] 3
##
$method
[1] "cosine"
##
$normalize_sim_matrix
[1] FALSE
##
$alpha
[1] 0.5
##
$verbose
[1] FALSE
```

(a) active user u8;

```
recsUBCF <- predict(modelUBCF, brm_u8, n=2)
recsUBCF

Recommendations as 'topNList' with n = 2 for 1 users.

getList(recsUBCF)

$u8
[1] "A" "B"

similarity(brm_offline,brm[8,],method="cosine")
```

```
u8
u1 0.5773503
u2 0.7071068
u3 0.0000000
u4 0.7071068
u5 0.0000000
u6 0.7071068
attr("method")
[1] "cosine"
attr("type")
[1] "simil"

recsIBCF <- predict(modelIBCF, brm_u8, n=2)
recsIBCF

Recommendations as 'topNList' with n = 2 for 1 users.

getList(recsIBCF)

$u8
[1] "A" "B"
```

(b) active user u7.

```
recsUBCF <- predict(modelUBCF, brm_u7, n=2)
getList(recsUBCF)

$u7
[1] "A" "B"

similarity(brm_offline, brm[7,], method="cosine")

u7
u1 0.4082483
u2 0.5000000
u3 0.3535534
u4 0.5000000
u5 0.3535534
u6 0.5000000
attr("method")
[1] "cosine"
attr("type")
[1] "simil"

recsIBCF <- predict(modelIBCF, brm_u7, n=2)
recsIBCF

Recommendations as 'topNList' with n = 2 for 1 users.

getList(recsIBCF)

$u7
[1] "G" "I"

similarity(brm[1:7,], method="cosine", which="items")

A B C G F I
B 0.3333333
C 0.7745967 0.5163978
G 0.0000000 0.4082483 0.0000000
F 0.0000000 0.3333333 0.2581989 0.8164966
I 0.0000000 0.4082483 0.0000000 1.0000000 0.8164966
J 0.0000000 0.0000000 0.0000000 0.7071068 0.5773503 0.7071068
```

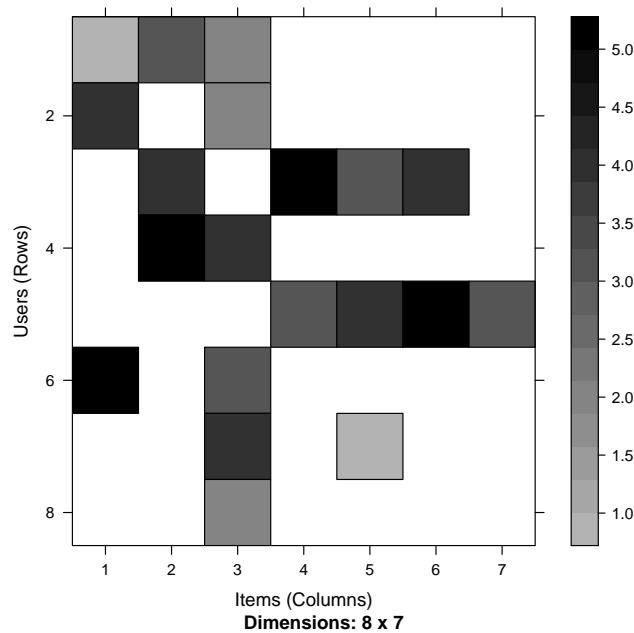
## Non-Binary Rating Data

13. Explore the types of recommendation models available for real rating matrices.

```
recommenderRegistry$get_entries(dataType = "realRatingMatrix")
```

14. Read the file `log1Ratings.csv`, containing information on the ratings given to web pages by a set of users, into a data frame in R. Build and deploy the following collaborative filtering recommendation models using, again, the first 6 users for training:

```
logR <- read_csv("log1Ratings.csv", col_types = list(col_factor(), col_factor(), col_integer()))
rrm <- as(as.data.frame(logR), "realRatingMatrix")
image(rrm)
```



```
rrm_offline <- rrm[1:6,]
getRatingMatrix(rrm_offline)

6 x 7 sparse Matrix of class "dgCMatrix"
A B C G F I J
u1 1 3 2
u2 4 . 2
u3 . 4 . 5 3 4 .
u4 . 5 4
u5 . . . 3 4 5 3
u6 5 . 3

similarity(rrm_offline, method="cosine")

u1 u2 u3 u4 u5
u2 0.8000000
u3 1.0000000 NA
u4 0.9962406 1.0000000 1.0000000
u5 NA NA 0.9400000 NA
u6 0.8436615 0.9970545 NA 1.0000000 NA

similarity(rrm_offline, method="cosine", which="items")

A B C G F I
B 1.0000000
C 0.9356015 0.9970545
G NA 1.0000000 NA
F NA 1.0000000 NA 0.9260924
I NA 1.0000000 NA 0.9374253 0.9995121
J NA NA NA 1.0000000 1.0000000 1.0000000
```

- (a) an user-based CF approach with two neighbours to predict the ratings of users u7 and u8;

```
modelUBCF_R <- Recommender(rrm_offline, "UBCF",parameter=list(nn=2))
getModel(modelUBCF_R)

$description
[1] "UBCF-Real data: contains full or sample of data set"
##
$data
6 x 7 rating matrix of class 'realRatingMatrix' with 17 ratings.
Normalized using center on rows.
##
$method
[1] "cosine"
##
$nn
[1] 2
##
$sample
[1] FALSE
##
$normalize
[1] "center"
##
$verbose
[1] FALSE

user u8
recsUBCF_R <- predict(modelUBCF_R,rrm[8,],type="ratings")
getList(recsUBCF_R)

$u8
A B G F I J
1 3 2 2 2 2

user u7
recsUBCF_R <- predict(modelUBCF_R,rrm[7,],type="ratings")
getList(recsUBCF_R)

$u7
A B G I J
3.0 2.5 3.0 2.5 2.5
```

- (b) an item-based CF approach with two neighbours to predict the ratings of users u7 and u8.

```
modelIBCF_R <- Recommender(rrm_offline, "IBCF",parameter=list(k=2))
getModel(modelIBCF_R)

$description
[1] "IBCF: Reduced similarity matrix"
##
$sim
7 x 7 sparse Matrix of class "dgCMatrix"
A B C G F I J
A . 1.0000000 0.8164966 . . .
B 1.0000000 1
C 0.8164966 0.4472136
G 0.9216354 . 1
F . . . 0.9216354 . . 1
I . 1.0000000 1
J 1.0000000 1 .
##
$k
[1] 2
##
$method
[1] "Cosine"
##
$normalize
[1] "center"
##
$normalize_sim_matrix
[1] FALSE
```

```
##
$alpha
[1] 0.5
##
$na_as_zero
[1] FALSE
##
$verbose
[1] FALSE

user u7
recsIBCF_R <- predict(modelIBCF_R,rrm[7,],type="ratings")
getList(recsIBCF_R)

$u7
A G J
4 1 1

user u8
recsIBCF_R <- predict(modelIBCF_R,rrm[8,],type="ratings")
getList(recsIBCF_R)

$u8
named numeric(0)
```

15. Considering the log1 binary data, evaluate different recommendation strategies.

- (a) Set the seed to 2021. Use the function `evaluationScheme` to define an evaluation scheme that splits the data into train and test set (80%-20% proportion) and establishes that 2 items of test cases are already known. In case that one or more users do not comply with this setting, you can disregard them.

```
log <- read_csv("log1.csv",col_types = list(col_factor(),col_factor()))
brm <- as(as.data.frame(log),"binaryRatingMatrix")

set.seed(2021) # for replication of results
e <- evaluationScheme(brm, method="split", train=0.8, given = 2)

Error in .local(data, ...): Some observations have size<given!

brm <- brm[rowCounts(brm)>=2,]
e <- evaluationScheme(brm, method="split", train=0.8, given = 2)
e

Evaluation scheme with 2 items given
Method: 'split' with 1 run(s).
Training set proportion: 0.800
Good ratings: NA
Data set: 7 x 7 rating matrix of class 'binaryRatingMatrix' with 19 ratings.
```

- (b) Check how the data was splitted according to the previous evaluation scheme, using the function `getData` on the evaluation scheme with the arguments "train", "known" and "unknown".

```
inspect(getRatingMatrix(brm))

items
[1] {A,B,C}
[2] {A,C}
[3] {B,G,F,I}
[4] {B,C}
[5] {G,F,I,J}
[6] {A,C}
[7] {C,F}

inspect(getRatingMatrix(getData(e,"train")))
```

```
items
[1] {C,F}
[2] {A,C}
[3] {A,C}
[4] {B,G,F,I}
[5] {G,F,I,J}

as(getRatingMatrix(getData(e,"train")), "matrix")

A B C G F I J
u7 FALSE FALSE TRUE FALSE TRUE FALSE FALSE
u6 TRUE FALSE TRUE FALSE FALSE FALSE FALSE
u2 TRUE FALSE TRUE FALSE FALSE FALSE FALSE
u3 FALSE TRUE FALSE TRUE TRUE TRUE FALSE
u5 FALSE FALSE FALSE TRUE TRUE TRUE TRUE

inspect(getRatingMatrix(getData(e,"known")))

items
[1] {A,B}
[2] {B,C}

inspect(getRatingMatrix(getData(e,"unknown")))

items
[1] {C}
[2] {}
```

- (c) Define the list of methods that will be used to obtain the top N recommendations, as follows:

```
methods <- list(
 "popular" = list(name="POPULAR", param = NULL),
 "user-based CF" = list(name="UBCF", param = NULL)
 "item-based CF" = list(name="IBCF", param = NULL)
)
```

- (d) Use the function `evaluate` with the previously defined evaluation scheme, methods and considering top 1, 3 and 5 recommendations for each of the models.

```
methods <- list(
 "popular" = list(name="POPULAR", param = NULL),
 "user-based CF" = list(name="UBCF", param = NULL),
 "item-based CF" = list(name="IBCF", param = NULL)
)

results <- evaluate(e, methods, type="topNList", n=c(1,3,5))

POPULAR run fold/sample [model time/prediction time]
1 [0.002sec/0.005sec]
UBCF run fold/sample [model time/prediction time]
1 [0sec/0.008sec]
IBCF run fold/sample [model time/prediction time]
1 [0.001sec/0.007sec]
```

- (e) Explore the obtained object.

```
results

List of evaluation results for 3 recommenders:
Evaluation results for 1 folds/samples using method 'POPULAR'.
Evaluation results for 1 folds/samples using method 'UBCF'.
Evaluation results for 1 folds/samples using method 'IBCF'.

class(results)
```

```
[1] "evaluationResultList"
attr(,"package")
[1] "recommenderlab"

avg(results) # just one run

$popular
TP FP FN TN precision recall TPR FPR
1 0.0 1.0 0.5 3.5 0.0000000 0 0 0.225
3 0.5 2.5 0.0 2.0 0.1666667 1 1 0.550
5 0.5 4.5 0.0 0.0 0.1000000 1 1 1.000
##
$`user-based CF`
TP FP FN TN precision recall TPR FPR
1 0.5 0.5 0 4 0.5000000 1 1 0.10
3 0.5 2.5 0 2 0.1666667 1 1 0.55
5 0.5 4.5 0 0 0.1000000 1 1 1.00
##
$`item-based CF`
TP FP FN TN precision recall TPR FPR
1 0.5 0.5 0 4 0.5000000 1 1 0.100
3 0.5 2.5 0 2 0.1666667 1 1 0.550
5 0.5 3.5 0 1 0.1250000 1 1 0.775

names(results)

[1] "popular" "user-based CF" "item-based CF"

results[["popular"]]

Evaluation results for 1 folds/samples using method 'POPULAR'.
```

- (f) Use the function `getConfusionMatrix` on one of the methods to obtain the corresponding confusion matrices. Be critical regarding the values that are shown.

```
getConfusionMatrix(results[["popular"]])

[[1]]
TP FP FN TN precision recall TPR FPR
1 0.0 1.0 0.5 3.5 0.0000000 0 0 0.225
3 0.5 2.5 0.0 2.0 0.1666667 1 1 0.550
5 0.5 4.5 0.0 0.0 0.1000000 1 1 1.000

model1 <- Recommender(getData(e,"train"), "POPULAR")
preds1 <- predict(model1,getData(e,"known"),n=3)
getList(preds1)

$u1
[1] "C" "F" "I"
##
$u4
[1] "F" "A" "I"

getConfusionMatrix(results[["user-based CF"]])

[[1]]
TP FP FN TN precision recall TPR FPR
1 0.5 0.5 0 4 0.5000000 1 1 0.10
3 0.5 2.5 0 2 0.1666667 1 1 0.55
5 0.5 4.5 0 0 0.1000000 1 1 1.00

model2 <- Recommender(getData(e,"train"), "UBCF")
preds2 <- predict(model2,getData(e,"known"),n=2)
getList(preds2)

$u1
[1] "C" "G"
##
$u4
[1] "A" "F"
```

```
getConfusionMatrix(results[["item-based CF"]])

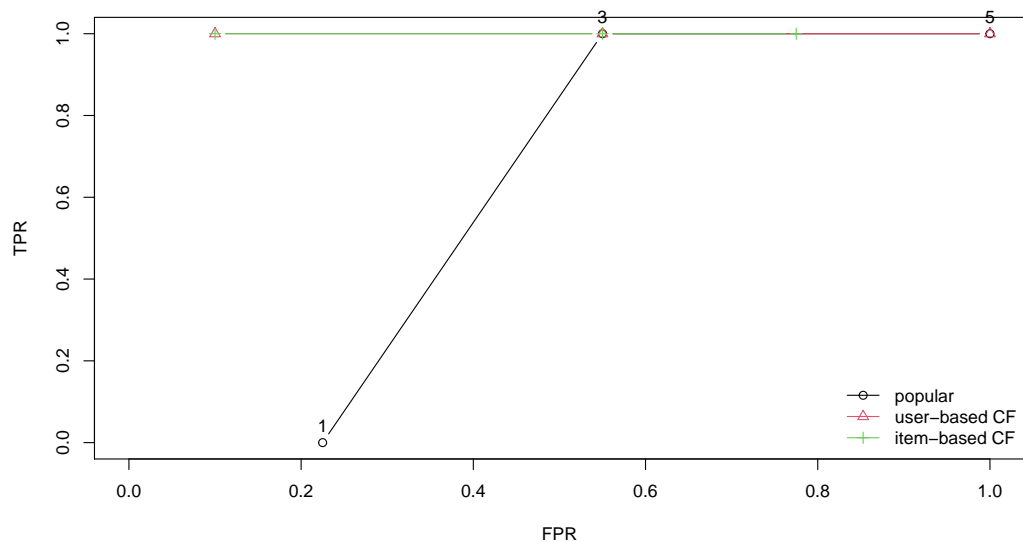
[[1]]
TP FP FN TN precision recall TPR FPR
1 0.5 0.5 0 4 0.5000000 1 1 0.100
3 0.5 2.5 0 2 0.1666667 1 1 0.550
5 0.5 3.5 0 1 0.1250000 1 1 0.775

model3 <- Recommender(getData(e,"train"), "IBCF")
preds3 <- predict(model3,getData(e,"known"),n=2)
getList(preds3)

$u1
[1] "C" "G"
##
$u4
[1] "A" "G"
```

(g) Plot the ROC curves for each of the methods and different values of N. What can you conclude?

```
plot(results, annotate=TRUE)
```



(h) Plot the precision/recall curves for each of the methods and different values of N. What can you conclude?

```
plot(results, "prec/rec", annotate=TRUE)
```



