Intelligent Data Mining - Exercise 4

Michael Debono Mrđen

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1 Assignment 1: Minhashing

- a. Compute the minhash signature for each column if we use the following three hash functions:
 - $h_1(x) = 2x + 1 \mod 6$
 - $h_2(x) = 3x + 2 \mod 6$
 - $h_3(x) = 5x + 2 \mod 6$

	Element	S_1	S_2	S_3	S_4	h_1	h_2	h_3
-	0	0	1	0	1	1	2	2
	1	0	1	0	0	3	5	1
	2	1	0	0	1	5	2	0
	3	0	0	1	0	1	5	5
	4	0	0	1	1	3	2	4
	5	1	0	0	0	5	5	3

- b. Which of these hash functions are true permutations? Only h_3 is a true permutation as it defines different hashes for all available
- c. How close are the estimated Jaccard similarities (based on the minhashes) for the six pairs of columns to the true Jaccard similarities.

Signature matrix computation:

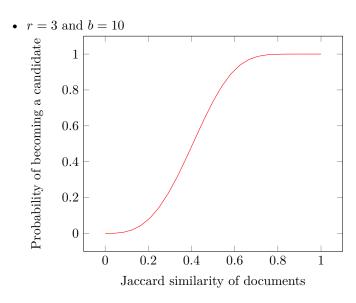
elements.

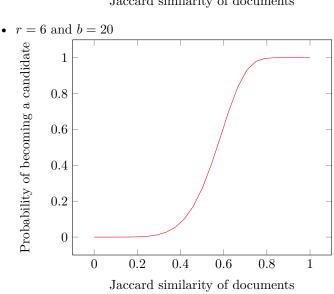
Jaccard similarities:

	Estimated	True	Difference
S_1, S_2	1/3	0	0.33
S_1, S_3	1/3	0	0.33
S_1, S_4	2/3	1/4	0.42
S_2, S_3	2/3	0	0.67
S_2, S_4	2/3	1/4	0.42
S_3, S_4	2/3	1/4	0.42

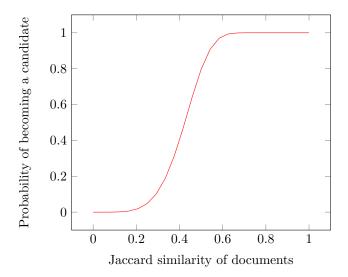
2 Assignment 2: Locality-sensitive hashing

a. Provide plots of the S-curve $1-(1-s^r)^b$ for the following values of r and b:





• r = 5 and b = 50



b. For each of the (r, b) pairs in (a), compute the threshold, that is, the value of s for which the value of $1 - (1 - s^r)^b$ is exactly 1/2. How does this value compare with the estimate of $(1/b)^{1/r}$ that was suggested in Section 3.4.2?

	s when $1 - (1 - s^r)^b = 1/2$	$(1/b)^{1/r}$	Difference
r = 3, b = 10	0.40609	0.46415	0.058
r = 6, b = 20	0.56935	0.60696	0.038
r = 5, b = 50	0.42439	0.45730	0.033

3 Assignment 3: Minhashing in Java

App.java:

```
public static long hash1(long itemID) {
            return (itemID + 1) % 9;
    public static long hash2(long itemID) {
            return (3*itemID + 1) % 9;
public static void main( String[] args ) throws IOException,
\hookrightarrow TasteException
{
    // load data
        DataModel data = new FileDataModel(new

→ File("data/data.csv"));
        // Represent each user as a set of item IDs (note:
        \rightarrow ignore ratings).
        // >> We will use the FastIDSet in the DataModel
        // Print the characteristic matrix (see Section
        \rightarrow 3.3.1) containing all users as columns and all
        \rightarrow item IDs as rows
        // (note: sort the matrix by Item ID).
        // Choose two hash functions (similar to Figure 3.4)
        → and compute the minhash signatures for the users
            // (similar to Section 3.3.5).
            // initialize user minhash signatures with max
             \rightarrow values
            HashMap<Long,Long> minh1s = new

→ HashMap<Long,Long>();
            HashMap<Long,Long> minh2s = new
             → HashMap<Long,Long>();
            LongPrimitiveIterator userIDsIterator =

    data.getUserIDs();

        while (userIDsIterator.hasNext()) {
                long userID = userIDsIterator.nextLong();
                minh1s.put(userID, Long.MAX_VALUE);
                minh2s.put(userID, Long.MAX_VALUE);
        }
        // Print characteristic matrix with hash signatures
        System.out.println("Characteristic matrix with hash

    signatures:");
            System.out.println();
```

{

```
// Print header line
System.out.print("
userIDsIterator = data.getUserIDs();
while (userIDsIterator.hasNext()) {
        long userID = userIDsIterator.nextLong();
        System.out.print(String.format(" | User %d",

    userID));
System.out.println(" || h_1 | h_2");
            System.out.println("-----
// Loop through all items
LongPrimitiveIterator itemIDsIterator =

    data.getItemIDs();

while (itemIDsIterator.hasNext()) {
        long itemID = itemIDsIterator.nextLong();
        System.out.print(String.format("Item %d ",
        → itemID));
        // Compute the hashes for the item
        long h1 = hash1(itemID);
        long h2 = hash2(itemID);
        // Loop through all users
        userIDsIterator = data.getUserIDs();
        while (userIDsIterator.hasNext()) {
            long userID = userIDsIterator.nextLong();
            // Get the items for the user
            FastIDSet itemIDs =

→ data.getItemIDsFromUser(userID);
            if (itemIDs.contains(itemID)) {
                    // If the user reviewed this
                    \rightarrow item, print 1
                    System.out.print(" | 1
                    // Store the hash value if it is
                    → less than what is already
                    \hookrightarrow stored
                    if (h1 < minh1s.get(userID)) {</pre>
                            minh1s.put(userID, h1);
                    if (h2 < minh2s.get(userID)) {</pre>
                            minh2s.put(userID, h2);
                    }
            } else {
                    // If the user did not review
                    → this item, print 0
                    System.out.print("| 0
```

```
}
        // Print the hash values
        System.out.print(String.format("||
                                              %d ",
        \rightarrow h1));
        System.out.print(String.format("|
        \rightarrow h2));
        System.out.println();
}
    System.out.println();
    // Print minhash signature matrix for users
    System.out.println("Minhash signature matrix for

    users:");

    System.out.println();
    System.out.print("
    // Print header line
userIDsIterator = data.getUserIDs();
while (userIDsIterator.hasNext()) {
        long userID = userIDsIterator.nextLong();
        System.out.print(String.format(" | User %d",

    userID));
}
System.out.println();
// Print data for hash function 1
System.out.print("h_1");
userIDsIterator = data.getUserIDs();
while (userIDsIterator.hasNext()) {
        long userID = userIDsIterator.nextLong();
        System.out.print(String.format(" | %d ",

→ minh1s.get(userID)));
System.out.println();
// Print data for hash function 2
System.out.print("h_2");
userIDsIterator = data.getUserIDs();
while (userIDsIterator.hasNext()) {
        long userID = userIDsIterator.nextLong();
        System.out.print(String.format(" | %d ",

→ minh2s.get(userID)));
System.out.println();
System.out.println();
// Print out the resulting pairwise similarities
\hookrightarrow based on the minhashes.
```

```
System.out.println("Estimated pairwise

    similarities:");
             System.out.println();
             // Loop over all users
             LongPrimitiveIterator i1 = data.getUserIDs();
             while (i1.hasNext()) {
                      long u1 = i1.nextLong();
                      // Loop again on all users for second user in
                       \hookrightarrow pair
                      LongPrimitiveIterator i2 = data.getUserIDs();
                      while (i2.hasNext()) {
                           long u2 = i2.nextLong();
                           // Take unique pairs of users, assuming
                           → they come in order
                           if (u1 < u2) {
                                    // Count similarities
                                    int sim = 0;
                                    if (minh1s.get(u1) ==
                                    \rightarrow minh1s.get(u2)) {
                                             sim++;
                                    }
                                    if (minh2s.get(u1) ==
                                    _{\hookrightarrow} \quad \texttt{minh2s.get}(\texttt{u2})) \ \{
                                             sim++;
                                    }
                                    // Divide similarity count by 2
                                    → for two hash functions
                                    System.out.println(String.format(

    "User %d, User %d = %.2f",

                                    \hookrightarrow u1, u2, sim/2. ));
                      }
             }
    }
}
```