Question 2: Iris Data Collection INTE1129 Assessment 1B The maximum marks for Question 2 is 4 Instructions Question 1: Fingerprint data collection textbox below: (1 mark) Question 2: Iris Data Collection Mated 0001_L_000 vs 0001_L_001 Question 3: Multimodal operation 0001_L_000 vs 0001_L_002 0001_L_000 vs 0001_L_003 Start Over 0001_L_000 vs 0001_L_004 0001_R_000 vs 0001_R_001 0001_R_000 vs 0001_R_002 0001_R_000 vs 0001_R_003 0001_R_000 vs 0001_R_004 0002_L_000 vs 0002_L_001 0002_L_000 vs 0002_L_002 0002_R_000 vs 0002_R_001 0002 R 000 vs 0002 R 002 0004_L_000 vs 0004_L_001 0004_L_000 vs 0004_L_002 0004_R_000 vs 0004_R_001 0004_R_000 vs 0004_R_002 0007_L_000 vs 0007_L_001 0007_R_000 vs 0007_R_001 0008_L_000 vs 0008_L_001 0008_R_000 vs 0008_R_001 Non Mated 0001_L_000 vs 0011_R_002 0001 L 000 vs 0011 L 001 0001_L_000 vs 0010_R_002 0001_L_000 vs 0010_R_001 0001_R_000 vs 0010_R_000 0001_R_000 vs 0010_L_002 0001_R_000 vs 0010_L_001 0001 R 000 vs 0010 L 000 0002_L_000 vs 0009_R_001 0002_L_000 vs 0009_R_000

Look at the Iris database - Retica_sample_iris_DB. It has iris captures from users numbered 0000 to 0011. Each user has Left and Right iris samples and the numbers of samples vary from 1 to 5. 2a. Use this database to create a list of 20 mated and 20 non mated iris comparisons. Write your list in the

0002_R_000 vs 0009_L_001 0002_R_000 vs 0009_L_000 0004_L_000 vs 0008_R_001 0004 L 000 vs 0008 R 000 0004_R_000 vs 0008_L_001 0004_R_000 vs 0008_L_000 0007_L_000 vs 0007_R_001 0007_R_000 vs 0000_L_000 0008_L_000 vs 0000_R_000 0008 R 000 vs 0001 L 001 Mated Comparisons data collection Open the IrisAlgorithmDemo application. Go to Tools on the menu bar and select Clear Database to empty the RAM of any existing images. Click Enrol and enrol the iris sample based on your mated comparison list. Click Identify and choose the iris sample you want to compare with from your list. Note the score on the bottom right panel of the application screen. 2b. Repeat this process and record all the 20 mated scores below:(0.5 marks) Use the format:

001 L 001 vs 001 L 002 = 791 002 R 001 vs 002 R 002 = 403 and so on.....

0007_R_000 vs 0007_R_001 = 427

0008_L_000 vs 0008_L_001 = 732

0008 R 000 vs 0008 R 001 = 777

Start Over

Code

0001_R_000

0001_R_000

0001 R 000

0001_R_000

0002 L 000

0002_L_000

mark)

Code

Start Over

comparisons_col1

<chr>

0001_L_000

0002 L 000

27

30 31

33

36

41

<chr>

0001_L_000

0001 L 000

0001_L_000

0001_L_000

0001_R_000

0001_R_000

0001_R_000

0001_R_000

0002 L 000

0002_L_000

<chr>

1-10 of 20 rows

0001_L_000

0001_L_000

0001_L_000

0001_L_000

0001_R_000

3.0

2.0

1.0

0.0

\$breaks

[1]

[14]

[27]

[40]

[53]

[66]

[79]

[92]

[105]

[118]

[131]

[144]

[157]

[170]

0.0

Previous Topic

0

0

65

130

195

260

325

390

455

520

585

650

715

780

845

5

70

135

200

265

330

395

460

525

590

655

720

785

850

10

75

140

205

270

335

400

465

530

595

660

725

790

855

15

80

145

210

275

340

405

470

535

600

665

730

795

860

20

85

150

215

280

345

410

475

540

605

670

735

800

865

25

90

155

220

285

350

415

480

545

610

675

740

805

870

30

95

160

225

290

355

420

485

550

615

680

745

810

875

35

100

165

230

295

360

425

490

555

620

685

750

815

880

40

105

170

235

300

365

430

495

560

625

690

755

820

885

45

110

175

240

305

370

435

500

565

630

695

760

825

890

50

115

180

245

310

375

440

505

570

635

700

765

830

895

55

120

185

250

315

380

445

510

575

640

705

770

835

900

60

125

190

255

320

385

450

515

580

645

710

775

840

905

Frequency

comparisons_col1

39 mated_hist

40 nonmated_hist

comparisons_col1

29 iris_nonmated_dataframe

32 #observe the score distribution

34 mated_scores<-as.numeric(iris_mated_dataframe[,3])</pre>

35 nonmated_scores<-as.numeric(iris_nonmated_dataframe[,3])</pre>

1-10 of 20 rows

1-10 of 20 rows

mated comparisons and the scores:(0.5 mark)

Mated 0001_L_000 vs 0001_L_001 = 791 0001 L 000 vs 0001 L 002 = 566

0001_L_000 vs 0001_L_003 = 746 0001_L_000 vs 0001_L_004 = 618 0001 R 000 vs 0001 R 001 = 589 0001_R_000 vs 0001_R_002 = 541 $0001_R_000 \text{ vs } 0001_R_003 = 612$

0001 R 000 vs 0001 R 004 = 638 0002_L_000 vs 0002_L_001 = 482 0002_L_000 vs 0002_L_002 = 522 0002 R 000 vs 0002 R 001 = 693

0002_R_000 vs 0002_R_002 = 737 0004_L_000 vs 0004_L_001 = 503 0004 L 000 vs 0004 L 002 = 551 0004_R_000 vs 0004_R_001 = 443 0004_R_000 vs 0004_R_002 = 881 0007 L 000 vs 0007 L 001 = 316

13 iris_mated_dataframe 14 15 comparisons_col1 comparisons_col2 comparisons_score <chr> <chr> <chr> 791 0001_L_000 0001_L_001 0001_L_000 0001_L_002 566 0001_L_000 0001_L_003 746 0001_L_000 0001_L_004 618

589

541

612

638

482

522

Previous 1 2 Next

► Run Code

2c. Use the codeblock below to create and display a dataframe called <code>iris_mated_dataframe</code> that has all the

3 rep("0007_R_000", times=1), rep("0008_L_000", times=1), rep("0008_R_000", times=1))

6 "0004_R_002", "0007_L_001", "0007_R_001", "0008_L_001", "0008_R_001")

0001_R_001

0001_R_002

0001_R_003

0001_R_004

0002_L_001

0002 L 002

The iris comparator will not give a non mated score for security reasons. We will simulate the non mated scores by

iris_nonmated_dataframe. The dataframe must have the non mated comparisons and generated scores (0.5)

3 rep("0007_R_000", times=1), rep("0008_L_000", times=1), rep("0008_R_000", times=1))

6 "0008_L_001", "0008_L_000", "0007_R_001", "0000_L_000", "0000_R_000", "0001_L_001")

1 comparisons_col1<- c(rep("0001_L_000", times =4), rep("0001_R_000", times =4), rep("0002_L_000", 2 rep("0002_R_000", times=2), rep("0004_L_000", times=2), rep("0004_R_000", times=2), rep("0007_L_0

4 comparisons_col2<- c("0011_R_002", "0011_L_001", "0010_R_002", "0010_R_001", "0010_R_000", "0010_ 5 "0010_L_001", "0010_L_000","0009_R_001", "0009_R_000", "0009_L_001","0009_L_000", "0008_R_001", "

generating them using a normal distribution. Choose the mean as 300 and standard deviation as 50.

2d. Use the code block below to generate 20 non mated scores and assign it to a variable called

Non mated comparisons data collection.

1 comparisons_col1<-c(rep("0001_L_000", times =4), rep("0001_R_000", times =4), rep("0002_L_000", t

2 rep("0002_R_000", times=2), rep("0004_L_000", times=2), rep("0004_R_000", times=2), rep("0007_L_0

4 comparisons_col2<-c("0001_L_001", "0001_L_002", "0001_L_003", "0001_L_004", "0001_R_001", "0001_F 5 "0001_R_003", "0001_R_004", "0002_L_001", "0002_L_002", "0002_R_001", "0002_R_002", "0004_L_001"

7 comparisons_score<-c(791, 566, 746, 618, 589, 541, 612, 638, 482, 522, 693, 737, 503, 551, 443, 8

12 iris_mated_dataframe<-as.data.frame(cbind(comparisons_col1,comparisons_col2,comparisons_score))

► Run Code

7 comparisons_score<-round(rnorm(20, mean = 300, sd=50),0) # change the number of generated outputs 9 iris_nonmated_dataframe<-as.data.frame(cbind(comparisons_col1,comparisons_col2,comparisons_score) 10 iris_nonmated_dataframe 11 12

comparisons_score

<chr>

308

297

Previous 1 2 Next

comparisons_col2

<chr>

0011_R_002

0009 R 000

1 # Copy the code for your mated and non mated database generation here.

8 "0004_R_002", "0007_L_001", "0007_R_001", "0008_L_001", "0008_R_001")

0001_L_000 0011_L_001 332 0010_R_002 0001_L_000 304 0001_L_000 0010_R_001 279 0010_R_000 0001_R_000 347 0001_R_000 0010_L_002 330 0001_R_000 0010_L_001 255 0001_R_000 0010_L_000 329 0009 R 001 365 0002_L_000

✓ Data analysis 2e. Use the code block below to generate a plot of the mated and non mated distributions.(0.5 mark) Code Start Over ▶ Run Code

14 iris_mated_dataframe<-as.data.frame(cbind(comparisons_col1,comparisons_col2,comparisons_score)) 15 iris_mated_dataframe

20 comparisons_col1<- c(rep("0001_L_000", times =4), rep("0001_R_000", times =4), rep("0002_L_000", 21 rep("0002_R_000", times=2), rep("0004_L_000", times=2), rep("0004_R_000", times=2), rep("0007_L_000", times=2)

23 comparisons_col2<- c("0011_R_002", "0011_L_001", "0010_R_002", "0010_R_001", "0010_R_000", "0010_ 24 "0010_L_001", "0010_L_000", "0009_R_001", "0009_R_000", "0009_L_001", "0009_L_000", "0008_R_001",

26 comparisons_score<-round(rnorm(20, mean = 300, sd=50),0) # change the number of generated outputs

28 iris_nonmated_dataframe<-as.data.frame(cbind(comparisons_col1,comparisons_col2,comparisons_score)

22 rep("0007_R_000", times=1), rep("0008_L_000", times=1), rep("0008_R_000", times=1))

25 "0008_L_001", "0008_L_000", "0007_R_001", "0000_L_000", "0000_R_000", "0001_L_001")

37 mated_hist<-hist(mated_scores, plot=TRUE, breaks = seq(from=0, to=1000, by=30))

comparisons_col2

<chr>

0001_L_001

0001_L_002

0001_L_003

0001_L_004

0001_R_001

0001_R_002

0001_R_003

0001_R_004

0002_L_001

0002_L_002

0011_R_002

0011_L_001

0010_R_002

0010_R_001

0010_R_000

<chr>

comparisons_col2

38 nonmated_hist<-hist(nonmated_scores, plot=TRUE, breaks = seq(from=0, to=1000, by=5))

3 comparisons_col1<-c(rep("0001_L_000", times =4), rep("0001_R_000", times =4), rep("0002_L_000", t 4 rep("0002_R_000", times=2), rep("0004_L_000", times=2), rep("0004_R_000", times=2), rep("0007_L_000", times=2)

6 comparisons_col2<-c("0001_L_001", "0001_L_002", "0001_L_003", "0001_L_004", "0001_R_001", "0001_F 7 "0001_R_003", "0001_R_004", "0002_L_001", "0002_L_002", "0002_R_001", "0002_R_002", "0004_L_001"

9 comparisons_score<-c(791, 566, 746, 618, 589, 541, 612, 638, 482, 522, 693, 737, 503, 551, 443, 8

5 rep("0007_R_000", times=1), rep("0008_L_000", times=1), rep("0008_R_000", times=1))

42 plot(mated_hist\$mids, mated_hist\$counts, type='l', lwd=2, col=3, xlab = "mated scores", ylab = "(43 lines(nonmated_hist\$mids, nonmated_hist\$counts, lwd=2, col=2) 44 title("Mated and Non Mated score distribution")

comparisons_score

<chr>

791

566

746

618

589

541

612

638

482

522

<chr>

351

298

287

273

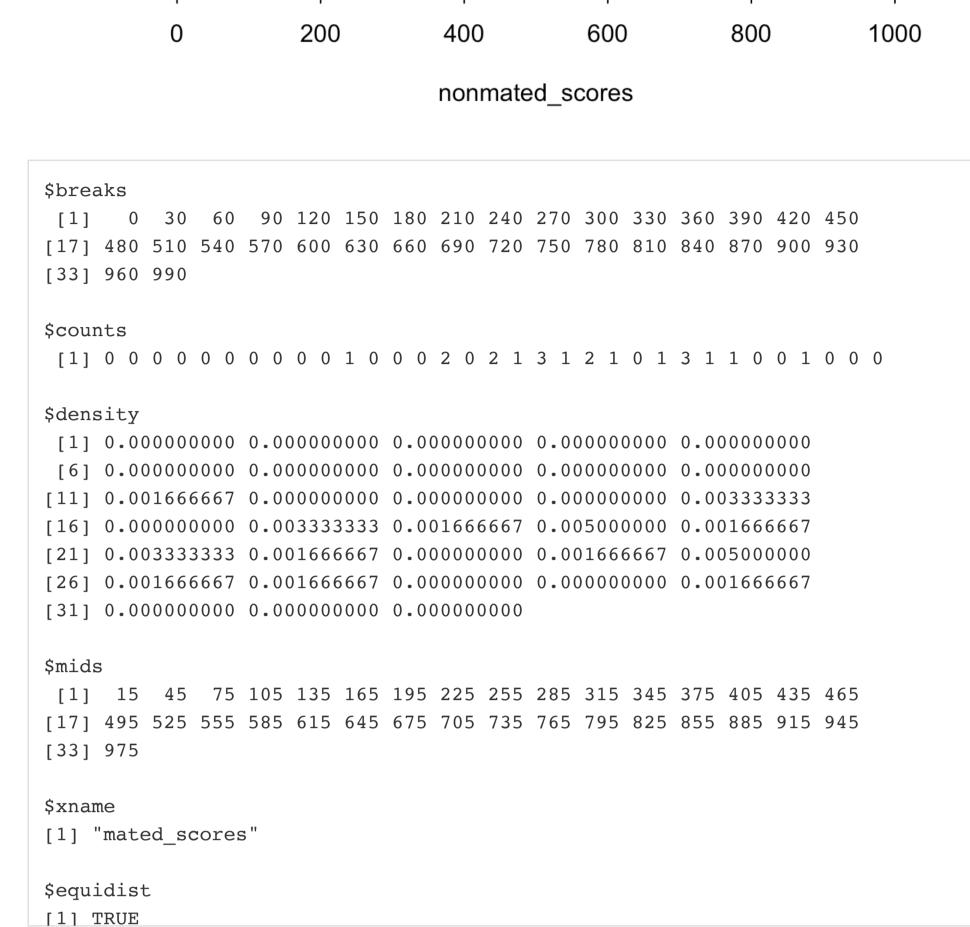
345

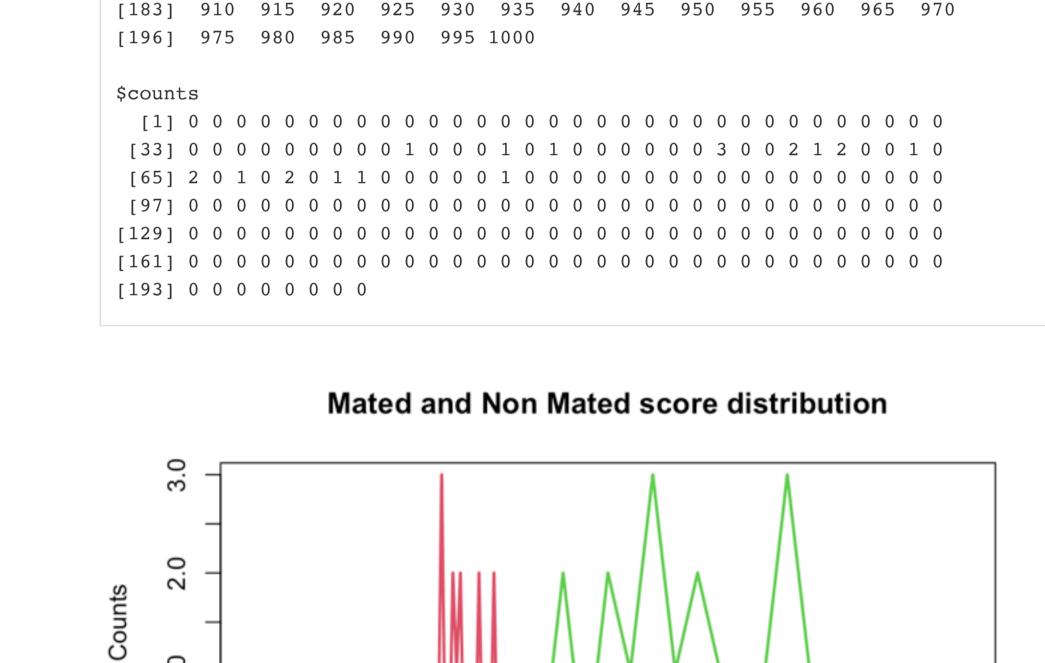
comparisons score

Previous 1 2 Next

0001_R_000 0010_L_002 236 0001_R_000 0010_L_001 275 0010_L_000 0001_R_000 230 0002_L_000 0009_R_001 207 0002_L_000 0009_R_000 299 Previous 1 1-10 of 20 rows 2 Next **Histogram of mated_scores** 3.0 2.0 Frequency 1.0 0.0 200 0 400 600 800 1000 mated_scores

Histogram of nonmated_scores





400

mated scores

2f. What would you choose as the operating threshold if this system was used to secure a high security

600

800

1000

none of the non-mated individuals can access the system and operating threshold of 450 is well above any sample for non-mated individuals.

Next Topic

clearance government document database? Justify your answer. (1 mark)

We would choose 450 as the threshold as the system being used is a high security

database. Having a higher threshold would eliminate the possibility of a non-mated

individual from accessing the system and accidental errors. We need to ensure that

200