

Quizzes due Apr 29, 2021 23:01 +03

Assignment Prep

Load the **GSE5859Subset** data:

```
library(GSE5859Subset)
data(GSE5859Subset)
```

Question 1

2/2 points (graded)

Inspect the dimensions of the **geneExpression** matrix.

A: How many samples are in the dataset?

✓ **Answer:** 8793

B: How many features are in the dataset?

✓ **Answer:** 24

Explanation

Part A:

```
dim(geneExpression)[1]
```

```
## [1] 8793
```

Part B:

```
dim(geneExpression)[2]
```

```
## [1] 24
```

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You have used 1 of 5 attempts

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Question 2

2/2 points (graded)

Inspect the `sampleInfo` data frame.

A: How many samples are from the ethnicity "ASN"?

23

✓ **Answer: 23**

23

B: Which sample is from the ethnicity "CEU"?

15

✓ **Answer: 15**

15

Explanation

Part A:

```
sum(sampleInfo$ethnicity == "ASN")
```

```
## [1] 23
```

Part B:

```
which(sampleInfo$ethnicity == "CEU")
```

```
## [1] 15
```

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Question 3

2/2 points (graded)

Inspect the `sampleInfo` data frame.

A: What is the distance between samples 3 and 7?

✓ **Answer:** 36.05656

B: What is the distance between samples 4 and 14?

✓ **Answer:** 46.72515

Explanation

Part A:

```
sqrt(crossprod(geneExpression[,3]-geneExpression[,7]))
```

```
##           [,1]
```

```
## [1,] 36.05656
```

Part B:

```
sqrt(crossprod(geneExpression[,4]-geneExpression[,14]))
```

```
##           [,1]
```

```
## [1,] 46.72515
```

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You have used 1 of 5
attempts

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Question 4

1/1 point (graded)

This code finds the mean distance between the first sample (column 1) and all other samples:

```
column = 1
x = 1:ncol(geneExpression)
dists = sapply(x, function(x){
  test = geneExpression[,x]
  target = geneExpression[,column]
  sqrt(crossprod(target-test))
})
mean(dists)
```

```
## [1] 34.2757
```

Add an extra `sapply()` loop to this code to check the mean distance between each sample (column) and all other samples.

Which sample (column) has the largest mean distance from other samples?
(Report the column number)

✓ **Answer: 15**

Explanation

```
columns = 1:ncol(geneExpression)
mean_dists = sapply(columns, function(column){
  dists = sapply(x, function(x){
    test = geneExpression[,x]
    target = geneExpression[,column]
    sqrt(crossprod(target-test))
  })
  mean(dists)
})
which.max(mean_dists)
```

```
## [1] 15
```

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You have used 1 of 5 attempts

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Question 5

1/1 point (graded)

Use `dist()` to calculate the distance between all pairs of samples.

What is the maximum distance between any two samples?

56.9792

✓ **Answer:** 56.97692

56.9792

Explanation

```
d = dist(t(geneExpression))  
max(d)
```

```
## [1] 56.97692
```

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You have used 2 of 5 attempts

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Question 6

2/2 points (graded)

A: What is the distance between features "1007_s_at" and "201371_s_at"?

12.42643

✓ **Answer:** 12.42643

12.42643

B: What is the distance between features "202138_x_at" and "202152_x_at"?

2.3093

✓ **Answer:** 2.3093

2.3093

Explanation

Part A:

```
sqrt(crossprod(geneExpression["1007_s_at",]-geneExpression[
```

```
##           [,1]
```

```
## [1,] 12.42643
```

Part B:

```
sqrt(crossprod(geneExpression["202138_x_at",]-geneExpression[
```

```
##           [,1]
```

```
## [1,] 2.3093
```

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You have used 1 of 5
attempts

i Answers are displayed within the problem

Question 7

1/1 point (graded)

Use `dist()` to calculate the distance between all pairs of features.

What is the maximum distance between any two features?

52.04894

✓ **Answer:** 52.04894

52.04894

Explanation

```
d2 = dist(geneExpression)
max(d2)
```

```
## [1] 52.04894
```

Submit

You have used 1 of 5
attempts

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