

RNAseq DCM Analysis LumA vs. LumB

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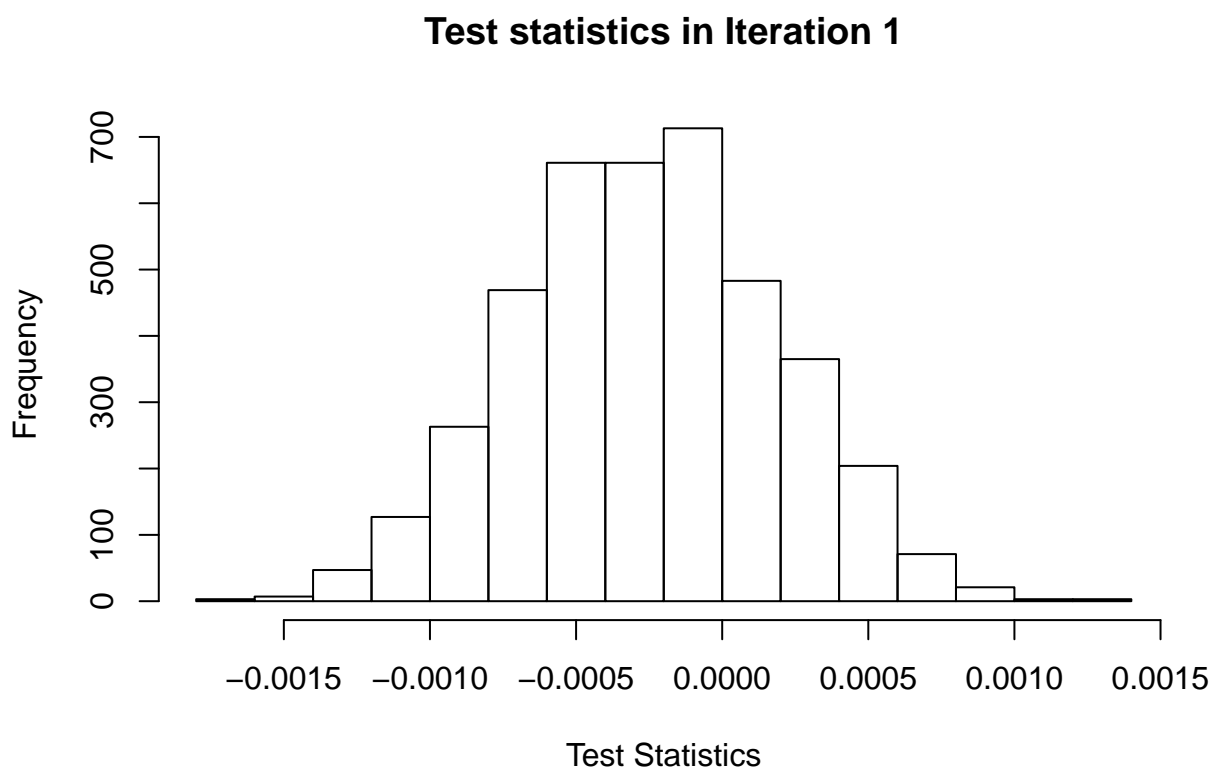
8/22/2018

This document includes data from a DCM run with RNAseq data after row-normalization.

$$\alpha = 0.05$$

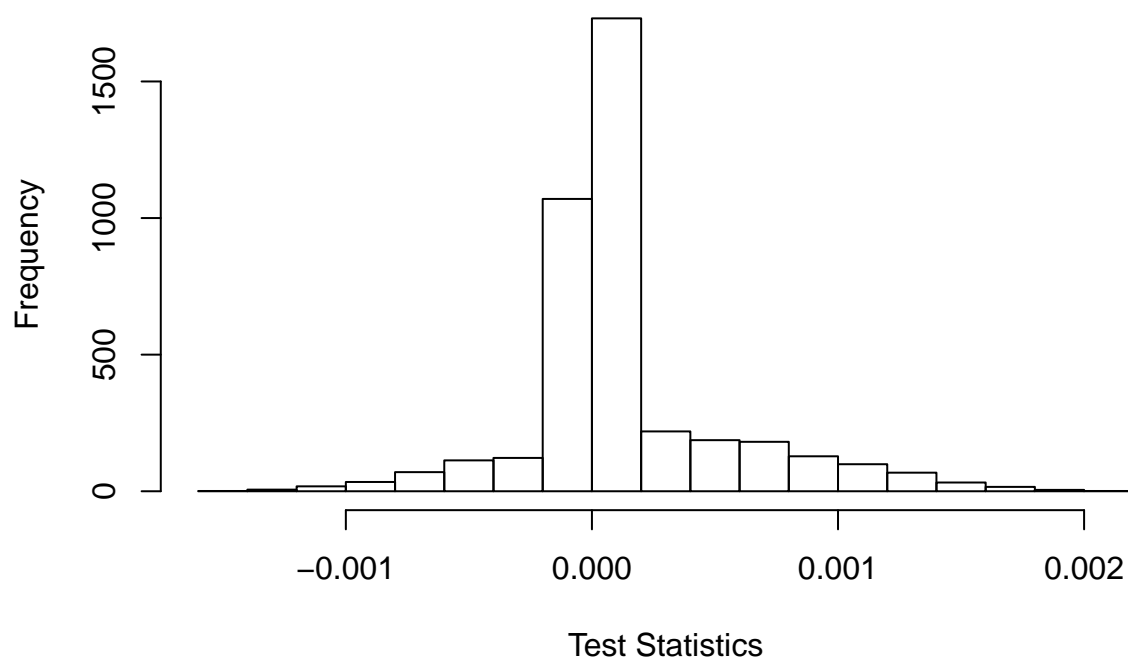
$QN = FALSE$

```
# Load data.  
load(file.path(data.dir, "DCM_1.RData"))  
  
# Plot test statistics.  
hist(DCM$it_test_stats[[1]], main="Test statistics in Iteration 1", xlab="Test Statistics")
```



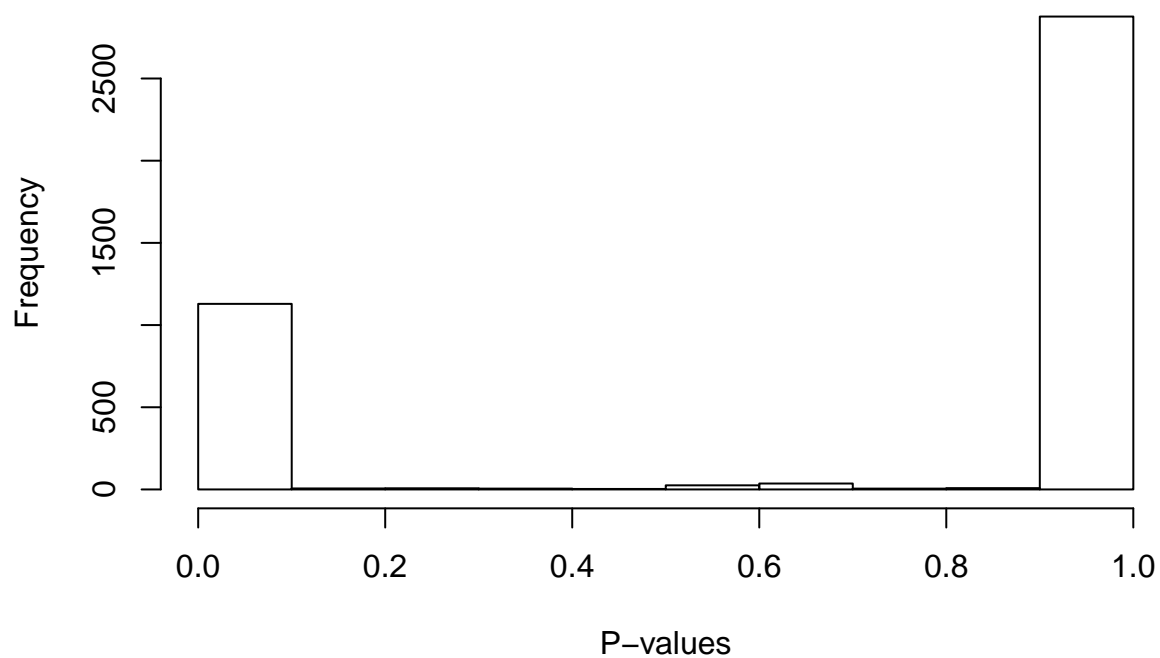
```
hist(DCM$it_test_stats[[11]], main="Test statistics in Iteration 11", xlab="Test Statistics")
```

Test statistics in Iteration 11



```
# Plot p-values.  
hist(DCM$it_p_vals[[1]], main="P-values in Iteration 1", xlab="P-values")
```

P-values in Iteration 1



```
hist(DCM$it_p_vals[[11]], main="P-values in Iteration 11", xlab="P-values")
```

P-values in Iteration 11

