

Sample number

90

60

30

0

0

5

10

15

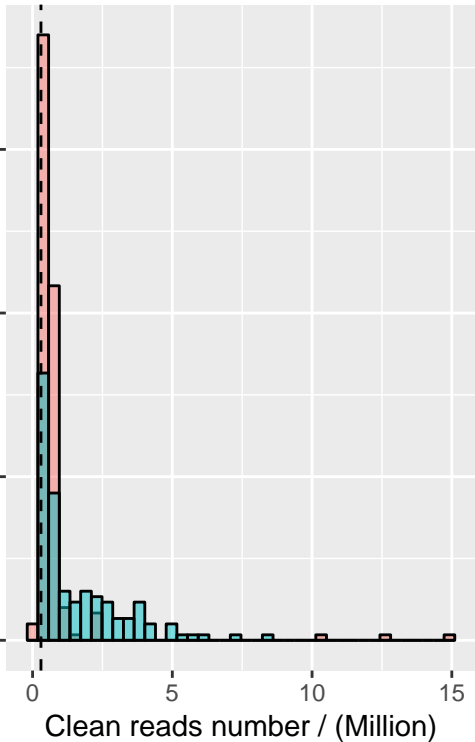
Clean reads number / (Million)

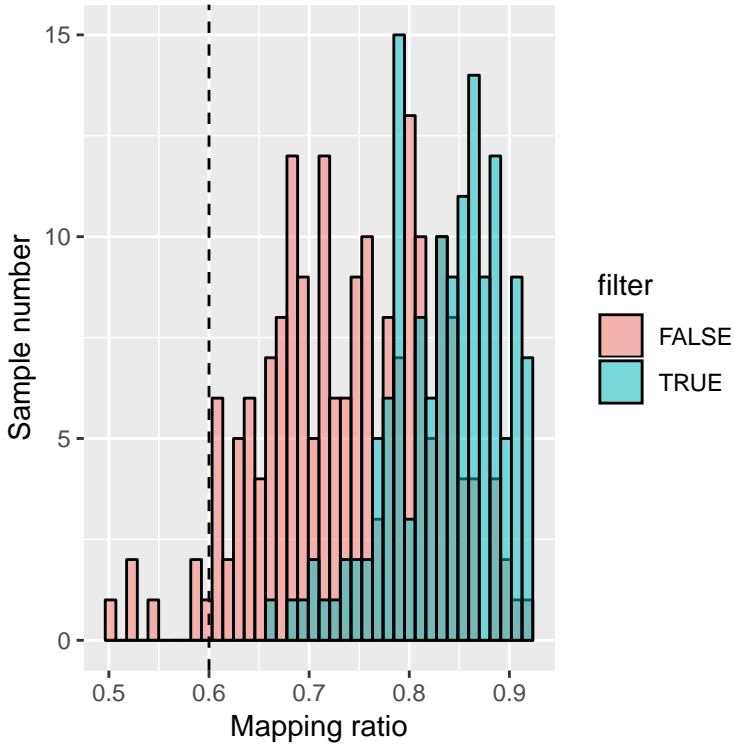
filter



FALSE

TRUE





Sample number

40

30

20

10

0

0

2500

5000

7500

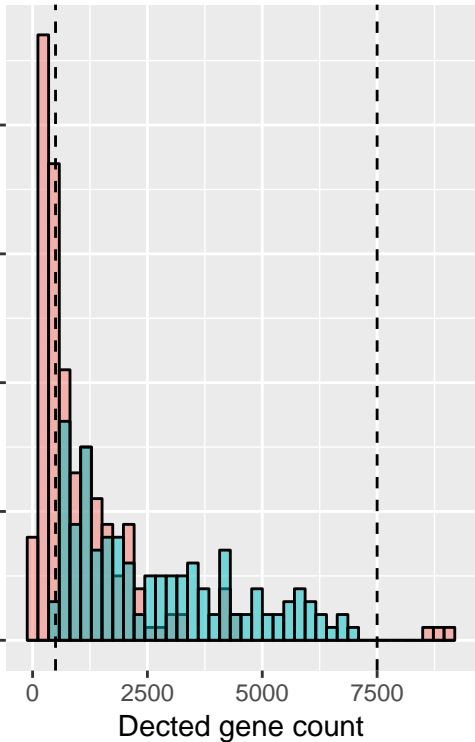
Dected gene count

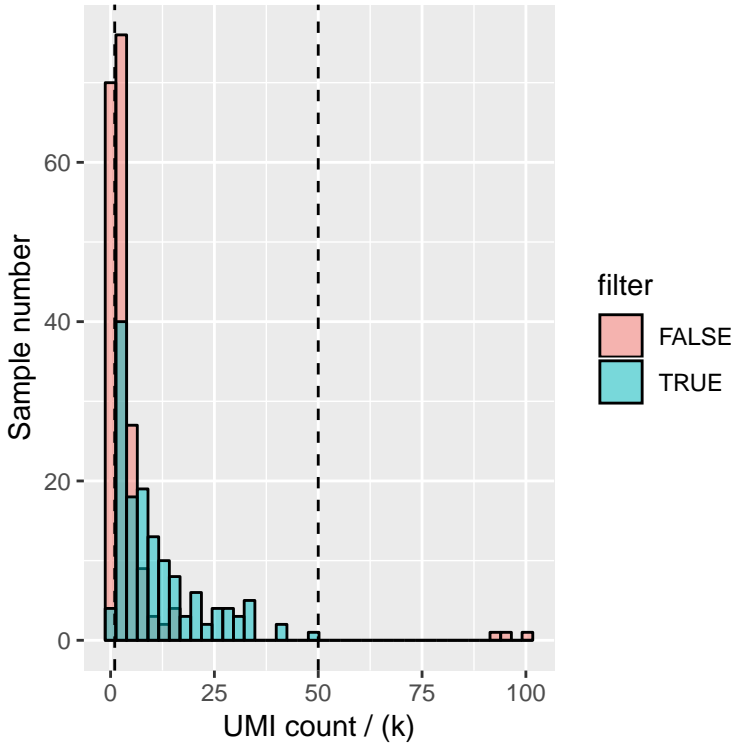
filter

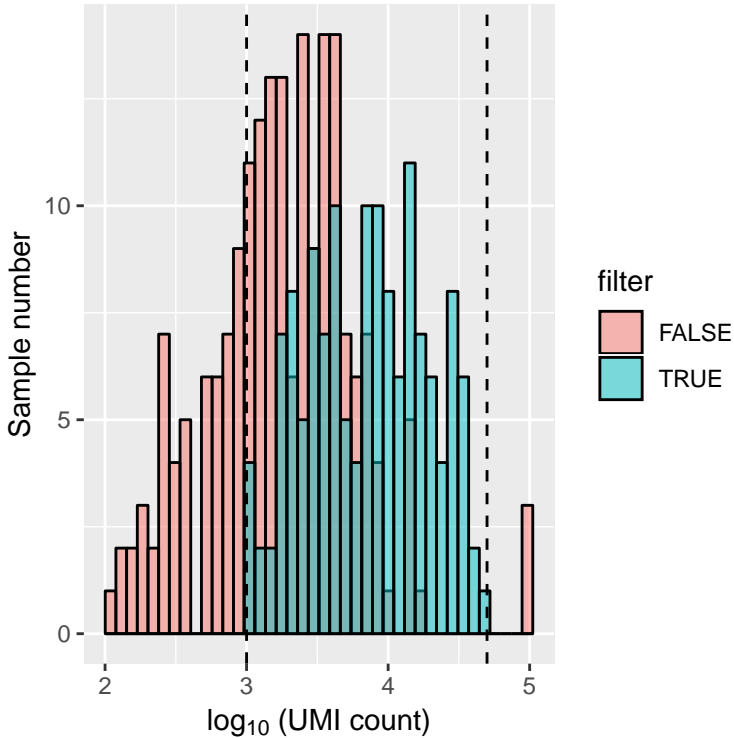


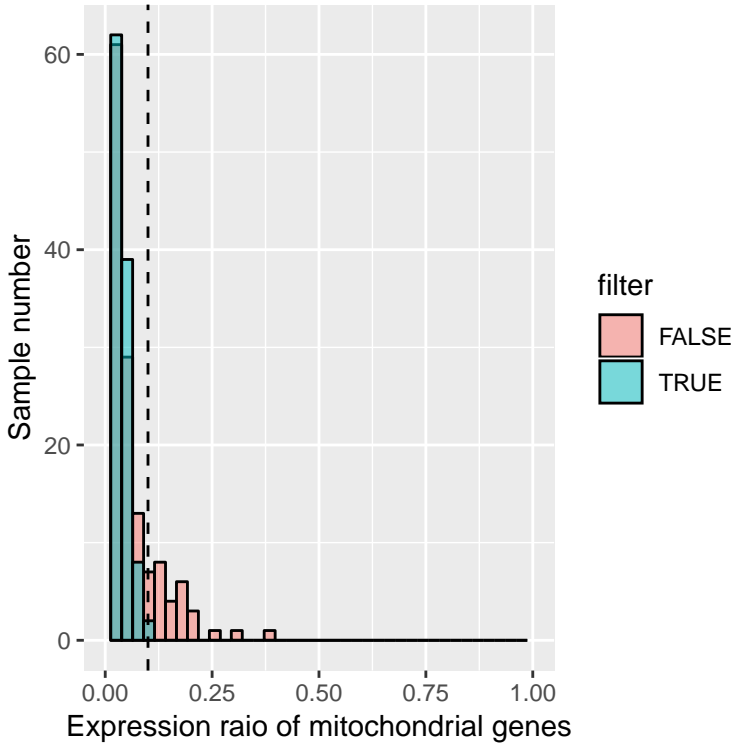
FALSE

TRUE

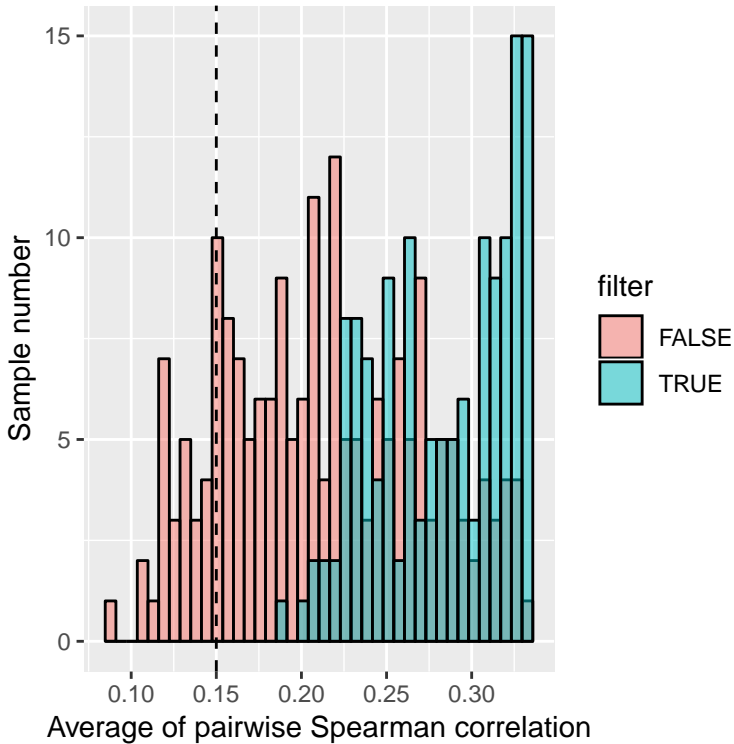












Average Spearman correlation

0.30  
0.25  
0.20  
0.15  
0.10

0

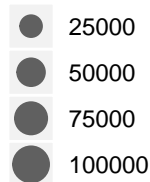
5

10

15

Reads number / (Million)

nUMI



filter



