

Number of detected genes

10000

8000

6000

0.0

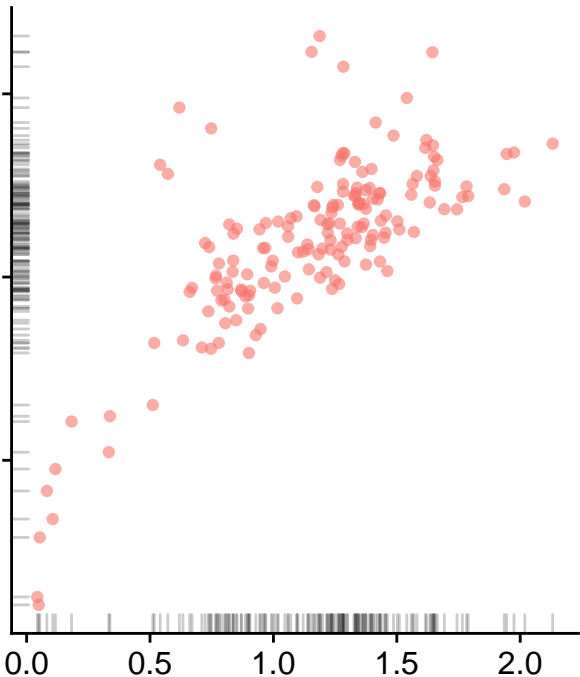
0.5

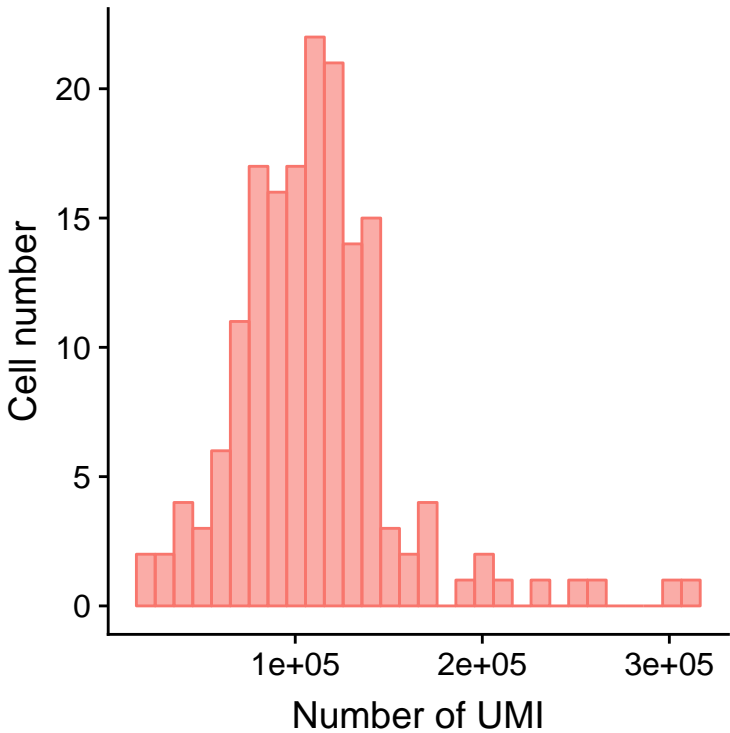
1.0

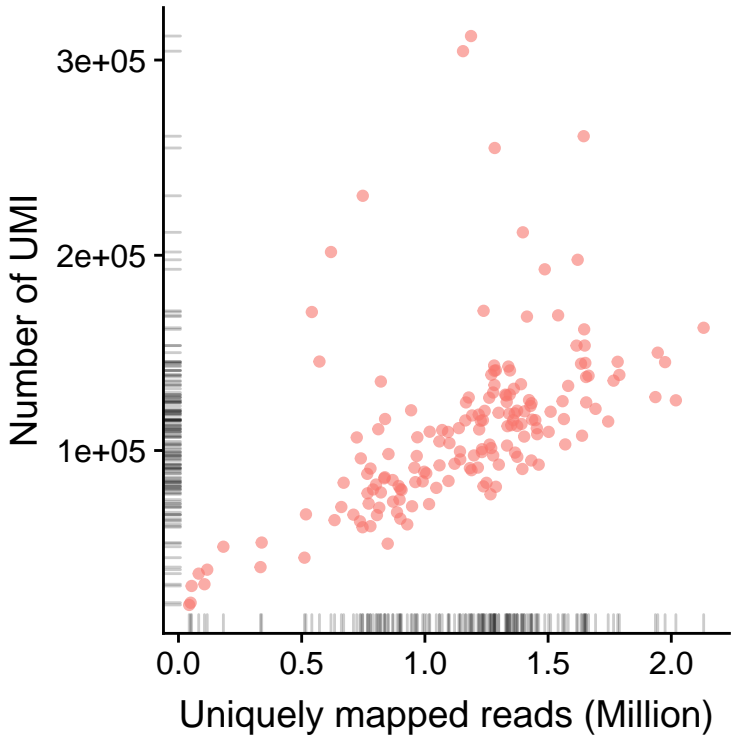
1.5

2.0

Uniquely mapped reads (Million)







Number of detected genes

10000

8000

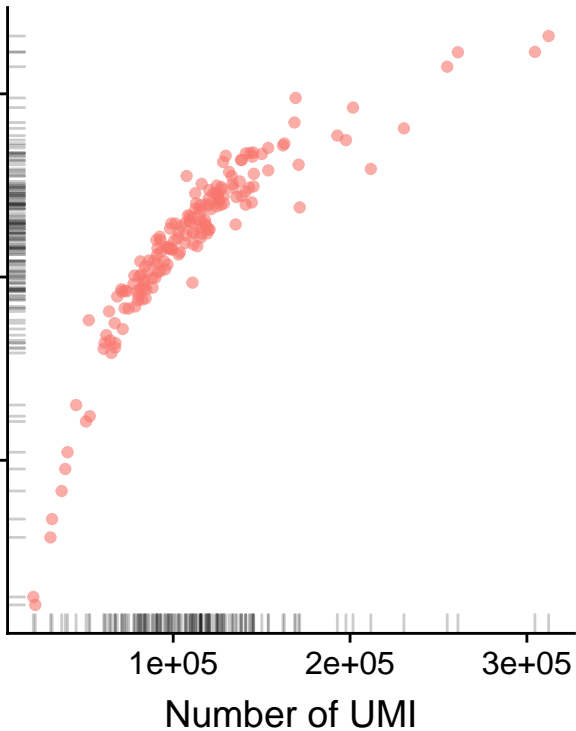
6000

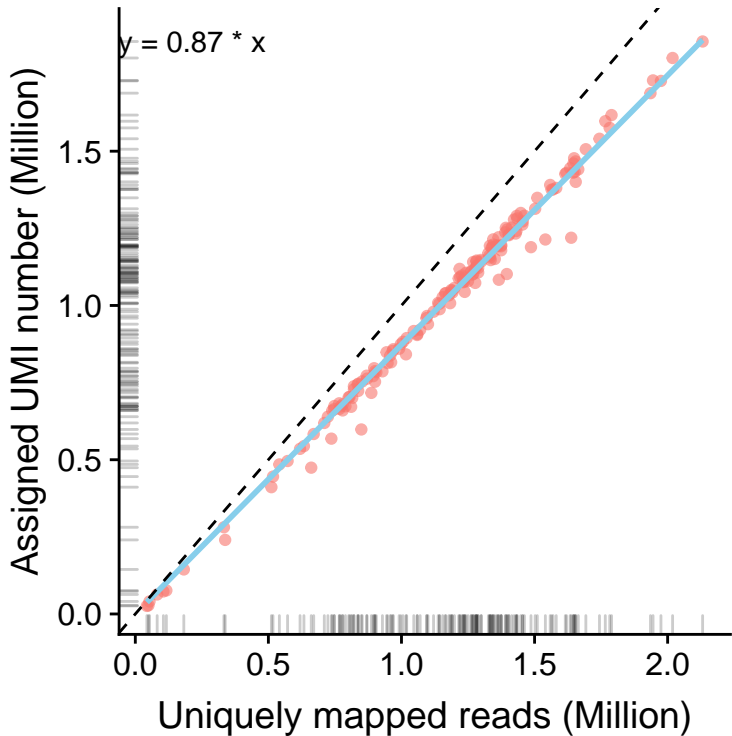
1e+05

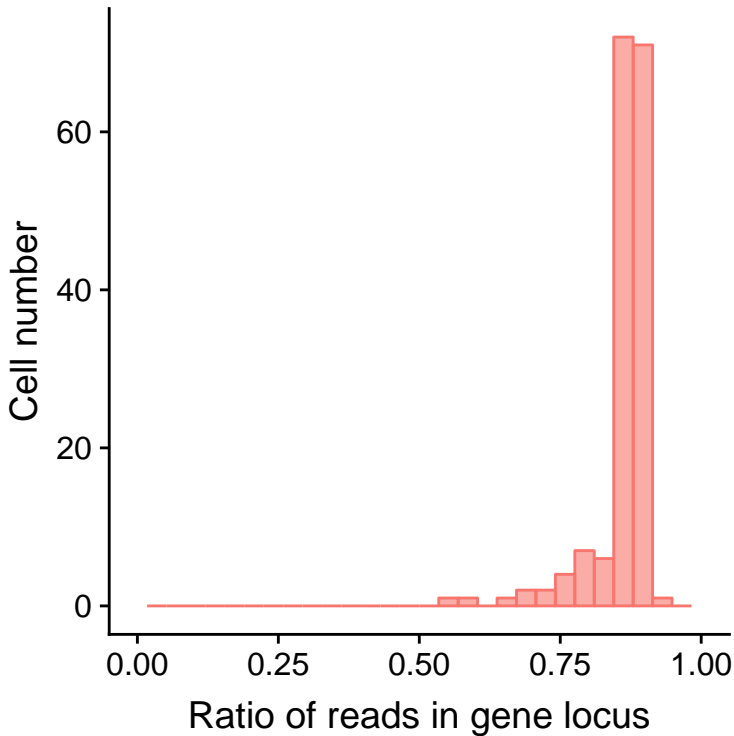
2e+05

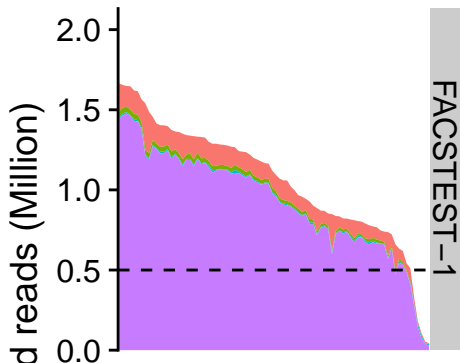
3e+05

Number of UMI

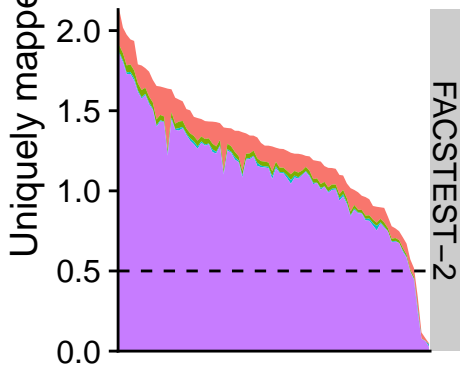




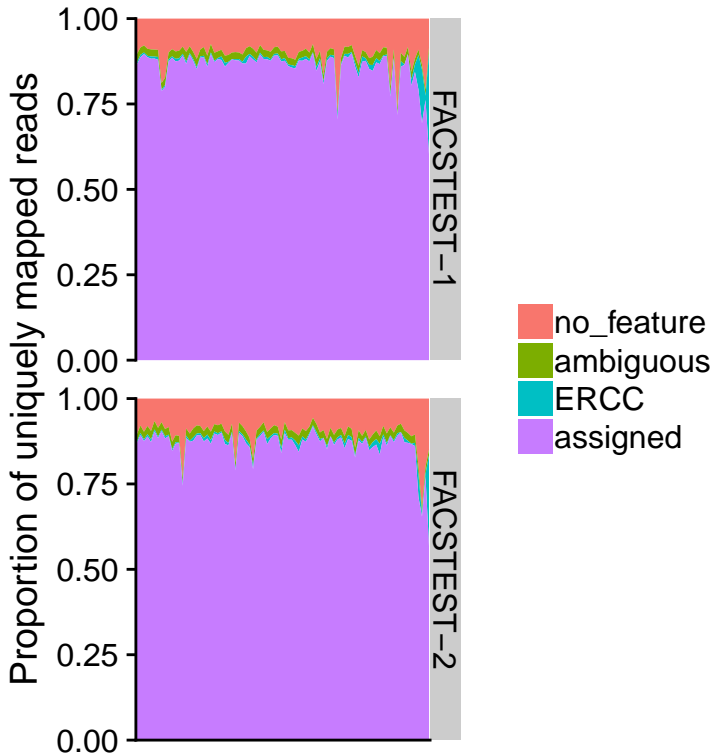


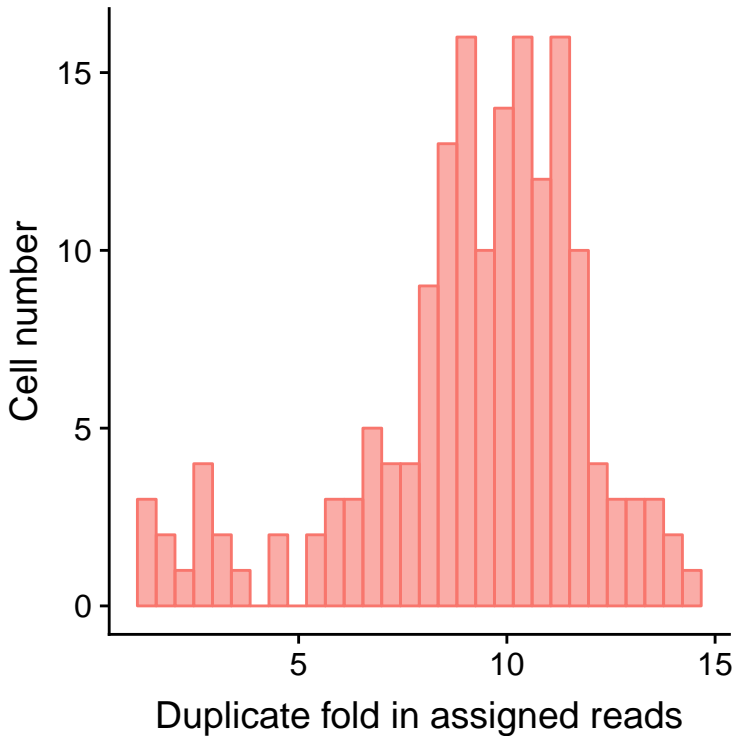


no\_feature  
ambiguous  
ERCC  
assigned









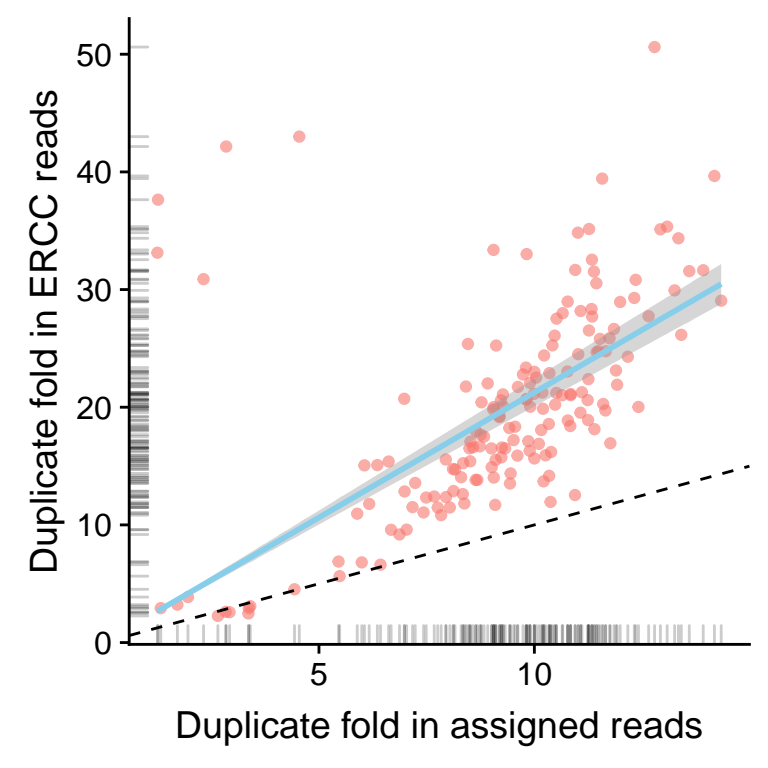
Duplicate fold in ERCC reads

50  
40  
30  
20  
10  
0

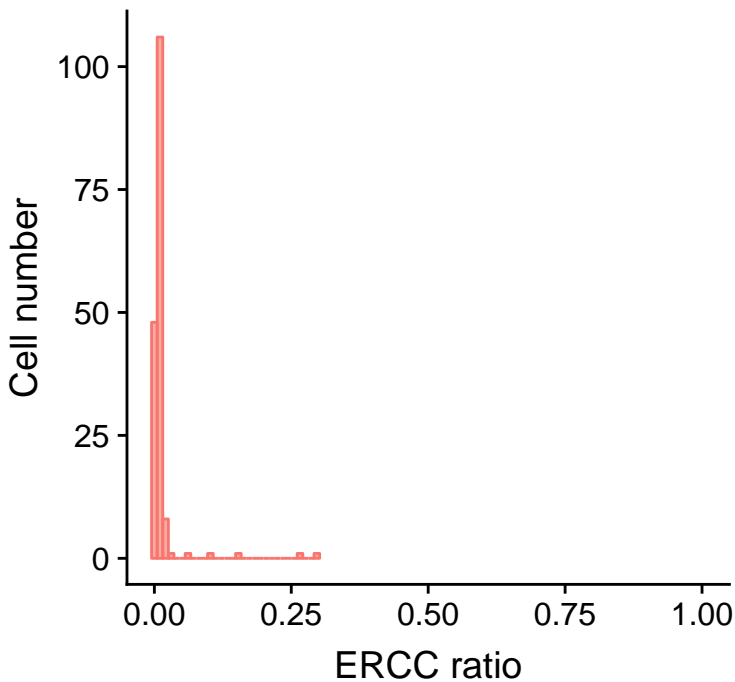
5

10

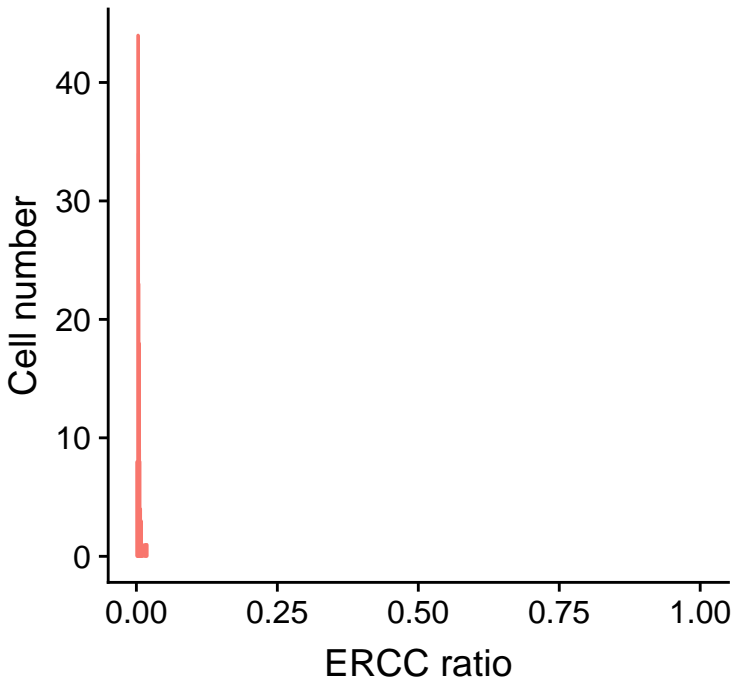
Duplicate fold in assigned reads



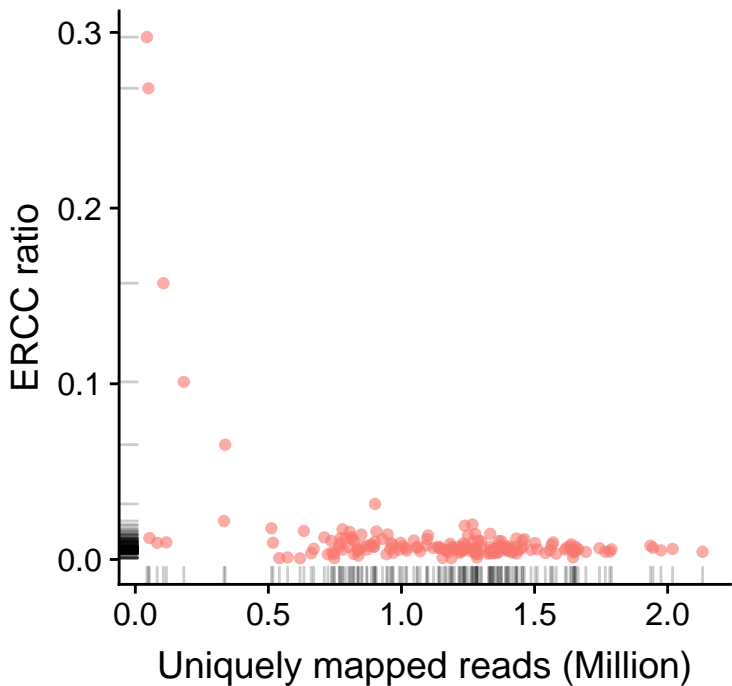
## Before removing duplicates



## After removing duplicates



## Before removing duplicates



## After removing duplicates

